

(19) World Intellectual Property
Organization
International Bureau



(43) International Publication Date
22 July 2004 (22.07.2004)

PCT

(10) International Publication Number
WO 2004/061410 A2

(51) International Patent Classification⁷: G01N
(21) International Application Number:
PCT/US2003/037090
(22) International Filing Date:
16 December 2003 (16.12.2003)
(25) Filing Language: English
(26) Publication Language: English
(30) Priority Data:
60/434,075 18 December 2002 (18.12.2002) US

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(81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.

(84) Designated States (regional): ARIPO patent (BW, GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),

[Continued on next page]

(54) Title: SERUM BIOMARKERS IN LUNG CANCER

MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION
IM-1	2011	A	IM-37	3893	A	IM-72	54028	A	IM-109	2882	B
IM-2	2030	A	IM-38	3960	A	IM-73	60170	A	IM-110	2967	B
IM-3	2069	A	IM-39	3972	A	IM-75	74372	A	IM-111	2977	B
IM-4	2128	A	IM-40	3984	A	IM-76	75545	A	IM-112	2994	B
IM-5	2146	A	IM-41	4066	A	IM-77	77543	A	IM-113	3031	B
IM-6	2186	A	IM-42	4178	A	IM-78	79507	A	IM-114	3048	B
IM-7	2232	A	IM-43	4287	A	IM-79	89854	A	IM-115	3148	B
IM-8	2277	A	IM-44	4297	A	IM-80	101831	A	IM-116	3166	B
IM-9	2285	A	IM-45	4309	A	IM-81	104301	A	IM-117	3283	B
IM-10	2318	A	IM-46	4484	A	IM-82	125160	A	IM-118	3308	B
IM-11	2411	A	IM-47	4849	A	IM-83	132976	A	IM-119	3332	B
IM-12	2434	A	IM-48	4798	A	IM-84	149099	A	IM-120	3432	B
IM-13	2467	A	IM-49	5104	A	IM-85	2016	B	IM-121	3450	B
IM-14	2482	A	IM-50	5918	A	IM-86	2029	B	IM-122	3561	B
IM-15	2498	A	IM-51	6122	A	IM-87	2144	B	IM-123	3616	B
IM-16	2565	A	IM-52	6192	A	IM-88	2130	B	IM-124	3714	B
IM-17	2574	A	IM-53	6462	A	IM-89	2168	B	IM-125	3730	B
IM-18	2586	A	IM-54	6680	A	IM-90	2184	B	IM-126	3834	B
IM-19	2605	A	IM-55	7788	A	IM-91	2200	B	IM-127	3899	B
IM-20	2722	A	IM-56	8145	A	IM-92	2284	B	IM-128	3969	B
IM-21	2746	A	IM-57	8954	A	IM-93	2299	B	IM-129	3986	B
IM-22	2788	A	IM-58	9312	A	IM-94	2314	B	IM-130	3997	B
IM-23	2866	A	IM-59	9449	A	IM-95	2314	B	IM-131	4013	B
IM-24	2871	A	IM-60	10272	A	IM-96	2428	B	IM-132	4181	B
IM-25	2984	A	IM-61	11663	A	IM-97	2451	B	IM-133	4297	B
IM-26	3030	A	IM-62	13378	A	IM-98	2486	B	IM-134	4311	B
IM-27	3144	A	IM-63	14698	A	IM-99	2483	B	IM-135	4465	B
IM-28	3243	A	IM-64	15190	A	IM-100	2565	B	IM-136	4484	B
IM-29	3273	A	IM-64	68758	A	IM-101	2583	B	IM-137	4579	B
IM-30	3290	A	IM-65	15951	A	IM-102	2597	B	IM-138	4608	B
IM-31	3369	A	IM-66	15172	A	IM-103	2697	B	IM-139	4669	B
IM-32	3445	A	IM-67	15925	A	IM-104	2715	B	IM-140	4747	B
IM-33	3483	A	IM-68	23436	A	IM-105	2740	B	IM-141	4862	B
IM-34	3676	A	IM-69	39794	A	IM-106	2752	B	IM-142	4891	B
IM-35	3779	A	IM-70	44166	A	IM-107	2767	B	IM-143	5033	B
IM-36	3793	A	IM-71	46890	A	IM-108	2865	B	IM-144	5077	B

(57) Abstract: Certain biomarkers and biomarker combinations are useful in a qualifying lung cancer status in a subject. A diagnostic methodology employing these biomarkers and combinations can detect whether a subject has lung cancer.

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European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

Published:

- *without international search report and to be republished upon receipt of that report*

SERUM BIOMARKERS IN LUNG CANCER

BACKGROUND OF THE INVENTION

[0001] The present invention relates generally to the field of serum biomarkers in lung carcinoma. More particularly, the invention relates to serum biomarkers that can distinguish lung cancer from normal.

[0002] Lung cancer is the leading cause of cancer death worldwide, resulting in 150,000 deaths per year in the United States. The mortality rate from lung cancer is greater than the combined mortality from breast, prostate and colorectal cancers. On the basis of morphology, lung cancer can be broadly classified into four main categories namely, adenocarcinoma, squamous cell carcinoma, large cell undifferentiated carcinoma and small cell carcinoma. In Hong Kong from 1990 to 1996, the proportions for adenocarcinoma, squamous cell carcinoma, large cell undifferentiated carcinoma and small cell carcinoma are 45.5%, 27.5%, 4.7% and 10.3% respectively. Both squamous cell carcinoma and small cell carcinoma are strongly associated with a smoking history.

[0003] Adenocarcinoma, squamous cell carcinoma, and large cell undifferentiated carcinoma are usually referred as "non-small cell carcinoma." They are relatively chemo-resistant, and hence the mainstay of treatment is surgery. By contrast, small cell carcinoma has a higher propensity for distant metastases and is mainly treated by chemotherapy.

[0004] Biopsy can be used to diagnose lung cancer, but it is an invasive procedure and, therefore, less than desirable. Other diagnostic methods for lung cancer include ultrasound and computed tomography (CT) scan.

[0005] It would be highly desirable to have a biomarker or combination of biomarkers capable of distinguishing between lung cancer and normal cells. In addition, a simple test could aid in tracking treatment progress and even identify molecular targets for therapy. The literature on lung cancer diagnosis has not disclosed heretofore such a biomarker or combination of biomarkers, however.

SUMMARY OF THE INVENTION

[0006] In accordance with the present invention, biomarkers and combinations of biomarkers are used to identify lung cancer. The method successfully distinguishes between lung cancer and normal states, and can be used to identify the particular type of lung cancer. In one embodiment, a method for qualifying lung carcinoma status in a subject (e.g., a patient) comprises analyzing a biological sample from the subject for one or more of the top 50 biomarkers as shown in Figure 2 or Figures 4A and 4B. Thus, to assess overall lung cancer risk versus normal, a biomarker is selected from the group consisting of

(A) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, IM-155, IM-157, IM-176, IM-445, IM-177, IM-440, IM-468, IM-438, IM-547, IM-359, IM-436, IM-106, IM-455, IM-444, IM-158, IM-265, IM-50, IM-159, IM-156, IM-439, IM-157, IM-508, IM-514, IM-478, IM-473, IM-360, IM-435, IM-150, IM-151, IM-110, IM-51, IM-163, IM-437, IM-546, IM-153, and IM-268, or

(B) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, WM-70, WM-369, WM-17, WM-473, WM-47, WM-203, WM-276, WM-279, WM-62, WM-366, WM-456, WM-428, WM-384, WM-287, WM-420, WM-292, WM-431, WM-455, WM-20, WM-340, WM-105, WM-389, WM-63, WM-354, WM-450, WM-466, WM-296, WM-343, WM-341, WM-339, WM-55, WM-66, WM-48, WM-38, WM-138, and WM-310,

[0007] wherein the biomarker is differentially present in samples of a subject with lung cancer and a so-called "normal" subject that is free of lung cancer.

[0008] More preferably, one or more of the top 15 biomarkers as shown in Figure 2 or Figures 4A and 4B is used to qualify lung cancer status. Thus, for assessing overall lung cancer status versus normal, the protein is selected from the group consisting of

(A) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, IM-155, IM-471, IM-510, IM-544, IM-474, and IM-155, or

(B) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, WM-70.

[0009] Still more preferably, one or more of the top 5 biomarkers as shown in Figure 2 or Figures 4A and 4B is used to qualify lung cancer status. In this instance, for overall lung cancer status versus normal, the biomarker is selected from the group consisting of

(A) IM-522, IM-273, IM-520, IM-519, and IM-454, or

(B) WM-61, WM-447, WM-446, WM-133, and WM-119.

[0010] In one embodiment, the method measures a plurality of biomarkers. The plurality of biomarkers can be measured simultaneously.

[0011] Biomarkers that, by themselves, are able to identify lung cancer include the WM-446 and WM-447 protein biomarkers, and these are particularly preferred.

[0012] The present invention also provides a method for qualifying lung cancer status in a subject (e.g., a patient), comprising (A) providing a spectrum generated by subjecting a biological sample from said subject to mass spectroscopic analysis that includes profiling on a chemically-derivatized affinity surface, and (B) putting the spectrum through pattern-recognition analysis that is keyed to at least one peak selected from the top 50 biomarkers as shown in Figure 2 or Figures 4A and 4B.

Thus, for qualifying overall lung cancer status, the biomarker is selected from the group consisting of

(i) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, IM-155, IM-157, IM-176, IM-445, IM-177, IM-440, IM-468, IM-438, IM-547, IM-359, IM-436, IM-106, IM-455, IM-444, IM-158, IM-265, IM-50, IM-159, IM-156, IM-439, IM-157, IM-508, IM-514, IM-478, IM-473, IM-360, IM-435, IM-150, IM-151, IM-110, IM-51, IM-163, IM-437, IM-546, IM-153, and IM-268 or

(B) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, WM-70, WM-369, WM-17, WM-473, WM-47, WM-203, WM-276, WM-279, WM-62, WM-366, WM-456, WM-428, WM-384, WM-287, WM-420, WM-292, WM-431, WM-455, WM-20, WM-340, WM-105, WM-389, WM-63, WM-354, WM-450, WM-466, WM-296, WM-343, WM-341, WM-339, WM-55, WM-66, WM-48, WM-38, WM-138, and WM-310.

[0013] For assessing the overall lung cancer status, the pattern-recognition analysis may, for example, be paired to a pair of peaks selected from the group consisting of (A) IM-266 and IM-474, IM-266 and IM-38, IM-266 and IM-454, IM-266 and IM-522, IM-266 and IM-544, IM-266 and IM-471, IM-474 and IM-151, IM-474 and IM-156, IM-474 and IM-544, IM-474 and IM-38, IM-522 and IM-507, IM-522 and IM-156, and IM-522 and IM-440;

or

(B) WM-447 and WM-59, WM-447 and WM-19, WM-447 and WM-118, WM-447 and WM-473, WM-19 and WM-59, WM-19 and WM-473, WM-19 and WM-369, WM-61 and WM-154, WM-61 and WM-369, WM-118 and WM-59 and WM-282 and WM-127.

[0014] More preferably, for assessing overall lung cancer status, the pattern-recognition analysis is keyed to a pair of peaks selected from the group consisting of (A) IM-266 and IM-474, IM-266 and IM-544, and IM-156 and IM-522;

or

(B) WM-447 and WM-59, WM-447 and WM-19, and WM-19 and WM-59.

[0015] Alternatively, the pattern-recognition analysis for assessing overall lung cancer status may be keyed to a triplet of peaks selected from the group consisting of

(A) IM-266, IM-454 and IM-474; and IM-266, IM-474 and IM-544;

or

(B) WM-447, WM-19 and WM-473.

[0016] In other embodiments, the pattern-recognition analysis may be keyed to a combination of more than three peaks, more particularly to a combination of 4, 5 or 6 peaks, where the combination is selected from among the combinations shown in Tables 1 and 2 herein.

[0017] In each case, the biomarker is differentially present in samples of a subject with lung cancer and a normal subject.

[0018] The invention also contemplates a kit for detecting and diagnosing lung cancer, thereby to assess lung cancer status. Kits within the invention comprise, for example, (i) an adsorbent attached to a substrate that retains one or more of the biomarkers shown in Figure 2 or Figures 4A and 4B, and (ii) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent. An inventive kit may further comprise a washing solution and/or instructions for making a washing solution. The kits may include more than type of adsorbent, each present on a different substrate, *e.g.*, on a WCX and IMAC biochip. In addition, the kits may comprise one or more containers with biomarker samples, to be used as standard(s) for calibration. The substrate comprising the adsorbent may be designed to engage a probe interface and, hence, function as a probe in gas phase ion spectrometry, preferably mass spectrometry. Alternatively, the kit may further comprise a second substrate adapted to engage the probe interface, on which the substrate comprising the adsorbent is mounted.

[0019] The method and kit according to the invention produce an article of manufacture in which one or more biomarkers according to the invention are bound to an adsorbent, optionally contacted with a matrix or energy absorbing molecule.

[0020] The present invention also provides software for qualifying lung carcinoma status in a subject, comprising an algorithm for analyzing data extracted from a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, wherein said data relates to one or more biomarkers according to the invention. In one embodiment, the algorithm carries out a pattern-recognition analysis that is keyed to data relating to at least one of the biomarkers. In another embodiment, the algorithm comprises classification tree analysis that is keyed to data relating to at least one of the biomarkers. In yet another embodiment, the algorithm

comprises an artificial neural network analysis that is keyed to data relating to at least one of the biomarkers.

[0021] In certain embodiments, the present invention provides methods and kits that use serum amyloid a protein or a fragment thereof to qualify lung carcinoma status in a subject. In one of these embodiments, the serum amyloid a biomarker has an apparent molecular weight of about 2803, 3168, 3277, 3552, 3897, 4300, 4490, 4655, 5927, 6874, 7776, 7941, 8152, 8952, 9233, 10300, 10866, or 10851 Daltons. In another embodiment, the serum amyloid a biomarker has an apparent molecular weight of about 3168, 3277, 3552, 3897, 4300, 4490, 4655, 7776, 7941, 8152, 8952, or 10851 Daltons. In yet another embodiment, the serum amyloid a biomarker has an apparent molecular weight of about 11.5 to 11.7 kD.

BRIEF DESCRIPTION OF THE DRAWINGS

[0022] Figures 1A-1D show all biomarkers identified with a Cu(II) IMAC3 ProteinChip® array format.

[0023] Figure 2 shows the top 50 biomarkers identified with a Cu(II) IMAC3 ProteinChip® array format.

[0024] Figures 3A-3O show all biomarkers identified with a WCX ProteinChip® array format.

[0025] Figures 4A and 4B show the top 50 biomarkers identified with a WCX ProteinChip® array format.

[0026] Figure 5 shows fragments of serum amyloid A (SAA) that are biomarkers according to the present invention.

[0027] Figure 6 shows identification of SAA biomarkers with an anti-SAA antibody.

[0028] Figures 7-16 are spectra from WCX chips in which all of the top 15 WCX marker peaks are labeled, along with various other peaks from among the top 50 WCX peaks. Red shows spectra from lung cancer patients and gray shows normals.

[0029] Figures 17-28 are spectra from IMAC chips in which all of the top 15 WCX marker peaks are labeled, along with various other peaks from among the top 50 IMAC peaks. Blue shows spectra from lung cancer patients and gray shows normals.

DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

[0030] In accordance with the present invention, a series of biomarkers associated with lung cancer has been discovered. In the present context, a biomarker is an organic biomolecule, particularly a polypeptide or protein, which is differentially present in a sample taken from a subject having lung cancer as compared to a comparable sample taken from a normal subject. A biomarker also may be differentially present in a sample taken from a subject with one type of lung cancer, *e.g.*, small cell carcinoma, as compared to a comparable sample taken from a subject with a different type of lung cancer, *e.g.*, adenocarcinoma or squamous cell carcinoma, or differentially present at different stages of a type of lung cancer. A biomarker is differentially present in samples taken from two groups of subjects if it is present at an elevated level or a decreased level in samples of the first group as compared to samples of the second group. More particularly, a biomarker is a polypeptide that is characterized by an apparent molecular weight, as determined by mass spectrometry, and that is present in samples from lung cancer subjects in an elevated or decreased level, as compared to subjects that do not have lung cancer. A biomarker is differentially present between two sets of samples if the amount of the biomarker in one sample set differs in a statistically significant way ($p < 0.01$) from the amount of biomarker in the other sample set.

[0031] The biomarkers of the invention can be used to assess lung cancer status in a subject. For example, they are capable of identifying lung cancer and successfully distinguishing it from normal subjects, thereby providing a way of diagnosing the presence or absence of lung cancer, including the presence or absence of a particular kind of lung cancer. In addition, the biomarkers are useful in assessing the risk of developing lung cancer, in staging of lung cancer and in assessing the effectiveness of treatment. Thus, "lung cancer status" in the context of the present invention includes, *inter alia*, the presence or absence of disease, the risk of developing disease, the stage of the disease, and the effectiveness of treatment of disease. Based on this status, further procedures may be indicated, including additional diagnostic tests or therapeutic procedures or regimens, such as endoscopy, biopsy, surgery, chemotherapy, immunotherapy, and radiation therapy.

[0032] In some instances, a single biomarker is capable of identifying lung cancer with a sensitivity or specificity of at least 85%, whereas, in other instances, a combination or plurality of biomarkers is used to obtain a sensitivity or specificity of at least 85%. The biomarkers and combinations of biomarkers thus can be used to qualify lung cancer status in a subject or patient.

[0033] The biomarkers according to the invention are present in serum. The biological sample used according to the present invention, however, need not be a serum sample. Thus, a biological sample for qualifying lung cancer status may be a serum, plasma or blood sample, although serum samples are preferred.

[0034] All of the biomarkers are characterized by molecular weight. A list of all the biomarkers obtained with the Cu(II) IMAC3 ProteinChip® array (CIPHERGEN Biosystems, Inc., Fremont, California, USA) is provided in Figures 1A-1D, and Figure 2 lists the top 50 biomarkers that distinguish between lung cancer and normal subjects that are identified by Cu(II) IMAC3 protocol described herein. Figures 3A-3O comprise a list of all the biomarkers obtained with the WCX2 ProteinChip® array, and Figures 4A and 4B comprise a ranking of the top 50 biomarkers that distinguish between (i) lung cancer and normal subjects, (ii) subjects with each of four types of lung cancer and normal subjects, and (iii) two types of lung cancer, *e.g.*, adenocarcinoma versus squamous cell carcinoma, as identified by WCX2 protocol described herein.

[0035] The top 50 biomarkers were determined by decision tree analysis using Biomarker Patterns™ software from CIPHERGEN Biosystems, Inc. Biomarkers other than those within the top 50 also are useful in distinguishing between subjects with lung cancer and normal subjects and may, in particular, appear in decision trees with multiple nodes. In preferred embodiments, one or more of the top 15 biomarkers are used, and in even more preferred embodiments, one or more of the top 5 biomarkers are used.

[0036] In each of Figures 1A-1D and 3A-3O, the number in the first column is the biomarker identifier. Thus, the first row in Figures 1A-1D relates to biomarker IM-1, the second row relates to biomarker IM-2, and so forth ("IM-" denoting biomarkers identified with the IMAC chip). Similarly, the first row in Figures 3A-3O relates to

biomarker WM-1 and the second row relates to biomarker WM-2 ("WM-" denoting biomarkers identified with the WCX2 chip). The number in the second column in Figures 1A-1D is the apparent molecular weight of the biomarker in daltons, as determined by mass spectrometry. In Figures 3A-3O, the apparent molecular weights for the biomarkers identified in the first column are reported in columns 3 through 11. The letter in the second column of Figures 1A-1D and the third column of Figures 3A-3O denotes the fraction in which the biomarker elutes in the protocol described herein; that is, biomarkers with an "A" elute in the first fraction, biomarkers with a "B" elute in the second fraction, and so forth. The fraction in which the biomarker elutes correlates with its pI, which biomarkers eluting at higher pH having a higher pI, and biomarkers eluting at lower pH having a lower pI.

[0037] Presenting the mass and affinity characteristics of a given biomarker within the invention, as in this description, characterizes that biomarker so as allow one to obtain and measure it, in accordance with the teachings herein. If desired, any of the biomarkers can be sequenced, in order to obtain an amino acid sequence, but this is not required to practice the present invention.

[0038] For example, a biomarker can be peptide mapped with a number of enzymes, such as trypsin and V8 protease, and the molecular weights of the digestion fragments can be used to search databases for sequences that match the molecular weights of the digestion fragments generated by the various enzymes. Alternatively, if the biomarkers are not proteins included in known databases, degenerate probes can be made based on the N-terminal amino acid sequence of the biomarker, which then are used to screen a genomic or cDNA library created from a sample from which the biomarker was initially detected. The positive clones can be identified, amplified, and their recombinant DNA sequences can be subcloned using techniques which are well known. Finally, protein biomarkers can be sequenced using protein ladder sequencing. Protein ladders can be generated by fragmenting the molecules and subjecting fragments to enzymatic digestion or other methods that sequentially remove a single amino acid from the end of the fragment. The ladder is then analyzed by mass spectrometry. The difference in masses of the ladder fragments identifies the amino acid removed from the end of the molecule.

[0039] Several biomarkers identified in accordance with the teachings of the present invention fit to serum amyloid A (SAA) or to a fragment of SAA. SAA is a well-known acute phase inflammatory marker. A number of the SAA biomarkers are identified in Figure 5 by both molecular mass and amino acid sequence. Most of these markers bound anti-SAA antibodies, as shown in Figure 6. The intact mass of SAA is 11.5 to 11.7 kD, and these biomarkers also have been identified by the present methodology. Fragments preferably have a molecular mass of at least about 200 Daltons, more preferably at least about 500 Daltons. In even more preferred embodiments, fragments have a molecular mass of at least about 800 Daltons, and most preferably at least about 1 Kilodalton.

[0040] In one embodiment, the fragments of SAA include a sequence of amino acids that is recognized by an epitope of an anti-SAA antibody. One way of identifying suitable fragments for use in the present invention is to enzymatically digest SAA and test the resulting fragments for the ability to bind to an anti-SAA antibody. Fragments that bind anti-SAA antibody can be sequenced using techniques well-known in the art, although the sequence of the fragment is not needed to practice the invention. In order to practice the invention with a fragment from the enzymatic digest that is identified as binding anti-SAA antibody, all that is required is to subject to the fragment to mass spectrometry to determine its mass.

[0041] The serum biomarkers according to the present invention were identified by comparing mass spectra of samples derived from sera from two groups of newly-diagnosed subjects, subjects with lung cancer and normal subjects. The subjects were diagnosed according to standard clinical criteria. Lung cancer subjects were histologically confirmed, and subjects without lung cancer were followed for at least 18 months following serum collection for any sign of lung cancer, to exclude subjects with asymptomatic lung cancer.

[0042] Sera from each group of subjects was collected, and fractionated with Q Ceramic HyperDF ion exchange resin (Biosepra SA, France) into six fractions which eluted at different pH. Fraction A comprised the flow through plus pH 9 eluant, Fraction B comprised the pH 7 eluant, Fraction C comprised the pH 5 eluant, Fraction D comprised the pH 4 eluant, Fraction E comprised the pH 3 eluant, and Fraction F

comprised isopropyl alcohol/acetonitrile TFA eluant. Fractions A through F are identified on Figures 7-28 as Fractions 1 through 6, respectively.

[0043] Each fraction was diluted and applied to a ProteinChip® array, either a Cu(II) IMAC3 or WCX2 chip array. Both of these chip arrays are produced by CIPHERGEN Biosystems, Inc. (Fremont, CA).

[0044] The Cu(II) IMAC3 is an "immobilized metal affinity-capture" chip, with a nitrilotriacetic acid surface for high-capacity copper binding and subsequent affinity capture of proteins with metal binding residues. Imidazole may be used in binding and washing solutions to moderate protein binding, including binding of non-specific proteins. Increasing the concentration of imidazole in the washing buffers reduces the binding of the target proteins. It is produced by photopolymerizing 5-methylacylamido-2-(N,N-bis(carboxymethylamino)pentanoic acid (7.5 wt%) and N,N'-methylenebisacrylamide (0.4 wt%) using (-) riboflavin (0.02 wt%) as a photoinitiator. The monomer solution is deposited onto the chip substrate and irradiated to photopolymerize. The chip then is activated with Cu(II).

[0045] The WCX2 is a weak cation exchange array with a carboxylate surface to bind cationic proteins. The negatively charged carboxylate groups on the surface of the WCX2 chip interact with the positive charges exposed on the target proteins. The binding of the target proteins is reduced by increasing the concentration of salt or by increasing the pH of the washing buffers.

[0046] Following application of the eluant fraction, the chips were incubated to allow the polypeptides in the eluant to bind to the sites on the chip by an affinity interaction. After incubation, each chip array was washed to remove polypeptides that bind non-specifically and buffer contaminants. That chip then was dried, and an energy absorbing molecule or matrix was applied to it, to facilitate desorption and ionization in a mass spectrometer.

[0047] In the mass spectrometer, retained polypeptides were desorbed from the chip array by laser desorption and ionization in a ProteinChip® Reader, which is integrated with ProteinChip® Software and a personal computer to analyze proteins captured on chip arrays. The ion optic and laser optic technologies in the ProteinChip® Reader detects proteins ranging from small peptides of less than 1000 Da up to proteins of

300 kilodaltons or more, and calculates the mass based on time-of-flight. Ionized polypeptides were detected and their mass accurately determined by this Time-of-Flight (TOF) Mass Spectrometry.

[0048] The mass spectra obtained for each group were subjected to scatter plot analysis, to eliminate run-to-run variation. Protein clusters on the scatter plot that had the same pattern for both lung cancer and normal subjects, *i.e.*, protein clusters that were either elevated in both groups of subjects or depressed in both groups of subjects, were eliminated as potential biomarkers. The remaining polypeptides were further analyzed for their ability to accurately identify subjects with lung cancer. Because the molecular weights were derived from scatter plot analysis, and because of limits on the ability of mass spectrometry to resolve molecular weights, the "absolute" molecular weight values given in Figures 1A-1D and 3A-3O actually represent approximate molecular weights.

[0049] The biomarkers of this invention are characterized by their mass-to-charge ratio as determined by mass spectrometry. The mass-to-charge ratio of each biomarker is provided in Figures 1A-1D and 3A-3O. For example, IM-1 in Figure 1A has a measured mass-to-charge ratio of 2011. The mass-to-charge ratios were determined from mass spectra generated on a Ciphergen Biosystems, Inc. PBS II mass spectrometer. This instrument has a mass accuracy of about +/- 0.15 percent. Additionally, the instrument has a mass resolution of about 400 to 1000 m/dm, where m is mass and dm is the mass spectral peak width at 0.5 peak height. The mass-to-charge ratio of the biomarkers was determined using Biomarker Wizard™ software (Ciphergen Biosystems). Biomarker Wizard assigns a mass-to-charge ratio to a biomarker by clustering the mass-to-charge ratios of the same peaks from all the spectra analyzed, as determined by the PBSII, taking the maximum and minimum mass-to-charge-ratio in the cluster, and dividing by two. Accordingly, the masses provided reflect these specifications.

[0050] The biomarkers of this invention are further characterized by the shape of their spectral peak in time-of-flight mass spectrometry. Mass spectra showing peaks representing the biomarkers are presented in Figures 7-28. The biomarker identifier numbers from Figures 2 and 4A-4B, respectively, are shown next to the peak, along

with their rank, which is indicated in parentheses below the biomarker identifier number.

[0051] The biomarkers of this invention are further characterized by their binding properties on chromatographic surfaces. Most of the biomarkers bind to IMAC (Cu) or WCX adsorbents (e.g., the CIPHERGEN® IMAC (Cu) or WCX ProteinChip® arrays) after washing as described herein.

[0052] Thus, a given molecular weight for a biomarker herein should be interpreted as the midpoint of a molecular-weight range. The accuracy of the mass spectrometer is $\pm 0.15\%$, and the actual molecular weight for a biomarker is therefore the value given, $\pm 0.15\%$. For example, the actual molecular weight for biomarker IM-273 is $11705 \pm 0.15\%$, or between 11687 and 11722. Often, the range surrounding the "absolute" value given in the figure is no more than ± 5 daltons (2006 to 2016 for IM-1), generally no more than ± 3 daltons (2008 to 2014 for IM-1), and often as small as ± 1 dalton (2010 to 2012 daltons for IM-1).

[0053] CART® (Salford Systems, San Diego, CA), a classification and regression tree software, was used to determine whether a potential biomarker had predictive value in assessing lung cancer. A software macro randomly selected a subset of 15% of the peaks from Figures 1A-1D or Figures 3A-3O. The peaks and peak heights from each sample were provided to the CART® software for analysis. The software performed an iterative analysis until a single decision tree was generated that was capable of distinguishing between cancerous and non-cancerous. Each node in the resulting decision tree sorted based on the peak height of a single biomarker. A tree may contain any number of nodes, but generally contains from 1 to 6 nodes. From a practical standpoint in a commercial diagnostic test, a decision tree with fewer nodes is preferred. A total of 2000 decision trees, each based on a different 15% subset of the peaks from Figures 1A-1D or Figures 3A-3O, were generated.

[0054] The CART® software assigned a score to each biomarker in the subset, based on its relative importance. A score of 100 is very high and a score of 0 is very low. The CART® software also determined the sensitivity and specificity of each decision tree.

[0055] The data generated by the decision tree analysis was subjected to further analysis. The biomarkers were ranked based on their average scores, which were determined by adding up a biomarker's scores for each decision tree in which it appeared, and dividing by the total number of decision trees in which the biomarker appeared. Approximately 500 of the potential biomarkers showed up in at least one tree, and most of the biomarkers showed up in about 150 to 400 of the two thousand trees. The top 50 biomarkers for the IMAC and WCX chip arrays as determined by this method are shown in Figures 2 and 4A-4B, respectively.

[0056] All of the trees having sensitivities and specificities greater than 85% also were identified. All trees capable of distinguishing lung cancer from normal and having from 1 to 6 nodes that meet the 85/85 criterion are shown in Tables 1 and 2.

TABLE 1. Decision trees with IMAC Biomarkers.

2 Nodes				
474	151			
474	156			
522	507		2 trees	
522	440		2 trees	
3 Nodes				
266	454	474		
474	156	153		
474	40	156		
520	276	113		
520	265	401		
522	151	474		
522	478	153		
522	156	474		
4 Nodes				
148	521	508	251	

266	544	474	493	
266	157	126	420	
266	544	474	482	
266	471	474	38	
266	544	474	38	
266	514	471	203	
522	58	266	474	
5 Nodes				
266	544	473	151	437
266	454	474	153	264
273	143	544	401	199

TABLE 2. Decision Trees with WCX Biomarkers.

1 Node					
446					
447					
2 Nodes					
282	127				
3 Nodes					
61	16	27			
61	119	154			
61	120	154			
61	369	184			
61	184	129			
61	19	282			
133	282	319			
282	59	218			
282	111	65			

446	19	16			
4 Nodes					
61	369	282	184		
61	48	203	3		
446	369	111	67		
446	466	58	120		
446	19	59	113		
446	282	19	47		
447	118	59	417		
447	118	59	473		
447	65	59	275		
447	19	59	282		
447	369	59	206		
447	19	59	253		
447	19	47	70		
5 Nodes					
61	369	128	184	197	
61	17	425	366	341	
133	139	363	216	273	
282	133	48	19	253	
369	310	19	109	384	
446	282	15	319	66	
447	19	71	473	31	
447	19	17	473	438	
447	47	31	365	59	
6 Nodes					
369	366	192	471	19	439

[0057] Each of the biomarker combinations of Tables 1 and 2 are preferred combinations for distinguishing lung cancer subjects from normal subjects in accordance with the present invention.

[0058] All biomarkers that appeared in at least two of the trees that met the 85/85 criterion were identified. For these biomarkers, Tables 3 and 4 provide the number of times the biomarker occurred in a trees that met the criterion, as well as the ranking of that biomarker on the top 50 lists of Figures 2 and 4A-4B.

TABLE 3. Correlation of IMAC biomarker decision tree frequencies and ranking.

Peak		# times		Rank
266		9		9
522		8		1
474		4		14
520		2		3
148		1		8
273		1		2

TABLE 4. Correlation of WCX biomarker decision tree frequencies and ranking.

Peak		# times		Rank
447		11		2
61		10		1
446		7		3
282		4		9
369		2		8
133		2		4

[0059] Biomarkers that occurred frequently in the highly discriminatory trees occurred among the top 50 ranked biomarkers, and typically had a top 10 ranking. In addition, certain pairs of biomarkers reappear, *e.g.*, WM-447 and WM-59, WM-447 and WM-19, WM-19 and WM-59, IM-266 and IM-474, IM-266 and IM-38, IM-266 and IM-454, IM-522 and IM-266. There also are repeats among triplets of biomarkers, such as IM-266, IM-266 and IM-38, and WM-447, WM-19 and WM-473. Other repeating pairs and trios of biomarkers can be seen in Tables 3 and 4, and are preferred.

[0060] Biomarkers and combinations of biomarkers identified in accordance with the present description may be used to qualify lung cancer status in a subject. In particular, a biomarker or combination of biomarkers can be used to distinguish lung cancer patients from normal patients with a high degree of specificity or sensitivity, *i.e.*, greater than at least 85%, preferably greater than at least 90%, and more preferably greater than 95%.

[0061] According to one aspect of the invention, therefore, the detection of biomarkers for diagnosis of lung cancer status entails contacting a sample from a subject with a substrate, *e.g.*, a SELDI probe, having an adsorbent thereon, under conditions that allow binding between the biomarker and the adsorbent, and then detecting the biomarker bound to the adsorbent by gas phase ion spectrometry, for example, mass spectrometry. Other detection paradigms that can be employed to this end include optical methods, electrochemical methods (voltametry and amperometry techniques), atomic force microscopy, and radio frequency methods, *e.g.*, multipolar resonance spectroscopy. Illustrative of optical methods, in addition to microscopy, both confocal and non-confocal, are detection of fluorescence, luminescence, chemiluminescence, absorbance, reflectance, transmittance, and birefringence or refractive index (*e.g.*, surface plasmon resonance, ellipsometry, a resonant mirror method, a grating coupler waveguide method or interferometry).

[0062] In one aspect, the markers of this invention are detect by gas phase ion spectrometry, which refers to the use of a gas phase ion spectrometer to detect gas phase ions. A gas phase ion spectrometer is an apparatus that detects gas phase ions. Gas phase ion spectrometers include an ion source that supplies gas phase ions. Gas

phase ion spectrometers include, for example, mass spectrometers, ion mobility spectrometers, and total ion current measuring devices.

[0063] "Mass spectrometer" refers to a gas phase ion spectrometer that measures a parameter which can be translated into mass-to-charge ratios of gas phase ions. Mass spectrometers generally include an ion source and a mass analyzer. Examples of mass spectrometers are time-of-flight, magnetic sector, quadrupole filter, ion trap, ion cyclotron resonance, electrostatic sector analyzer and hybrids of these. "Mass spectrometry" refers to the use of a mass spectrometer to detect gas phase ions. "Laser desorption mass spectrometer" refers to a mass spectrometer which uses laser as a means to desorb, volatilize, and ionize an analyte.

[0064] "Mass analyzer" refers to a sub-assembly of a mass spectrometer that comprises means for measuring a parameter which can be translated into mass-to-charge ratios of gas phase ions. In a time-of flight mass spectrometer the mass analyzer comprises an ion optic assembly, a flight tube and an ion detector.

[0065] "Ion source" refers to a sub-assembly of a gas phase ion spectrometer that provides gas phase ions. In one embodiment, the ion source provides ions through a desorption/ionization process. Such embodiments generally comprise a probe interface that positionally engages a probe in an interrogatable relationship to a source of ionizing energy (e.g., a laser desorption/ionization source) and in concurrent communication at atmospheric or subatmospheric pressure with a detector of a gas phase ion spectrometer.

[0066] Forms of ionizing energy for desorbing/ionizing an analyte from a solid phase include, for example: (1) laser energy; (2) fast atoms (used in fast atom bombardment); (3) high energy particles generated via beta decay of radionuclides (used in plasma desorption); and (4) primary ions generating secondary ions (used in secondary ion mass spectrometry). The preferred form of ionizing energy for solid phase analytes is a laser (used in laser desorption/ionization), in particular, nitrogen lasers, Nd-Yag lasers and other pulsed laser sources. "Fluence" refers to the laser energy delivered per unit area of interrogated image. Typically, a sample is placed on the surface of a probe, the probe is engaged with the probe interface and the probe

surface is struck with the ionizing energy. The energy desorbs analyte molecules from the surface into the gas phase and ionizes them.

[0067] Other forms of ionizing energy for analytes include, for example: (1) electrons which ionize gas phase neutrals; (2) strong electric field to induce ionization from gas phase, solid phase, or liquid phase neutrals; and (3) a source that applies a combination of ionization particles or electric fields with neutral chemicals to induce chemical ionization of solid phase, gas phase, and liquid phase neutrals.

[0068] A preferred mass spectrometric technique for use in the invention is Surface Enhanced Laser Desorption and Ionization (SELDI), as described, for example, in U.S. patents No. 5,719,060 and No. 6,225,047, both to Hutchens and Yip, in which the surface of a probe that presents the analyte (here, one or more of the biomarkers) to the energy source plays an active role in desorption/ionization of analyte molecules. In this context, "probe" refers to a device adapted to engage a probe interface and to present an analyte to ionizing energy for ionization and introduction into a gas phase ion spectrometer, such as a mass spectrometer. A probe typically includes a solid substrate, either flexible or rigid, that has a sample-presenting surface, on which an analyte is presented to the source of ionizing energy.

[0069] One version of SELDI, called "Surface-Enhanced Affinity Capture" or "SEAC," involves the use of probes comprised of a chemically selective surface ("SELDI probe"). A "chemically selective surface" is one to which is bound either the adsorbent, also called a "binding moiety" or "capture reagent," or a reactive moiety that is capable of binding a capture reagent, *e.g.*, through a reaction forming a covalent or coordinate covalent bond.

[0070] The phrase "reactive moiety" here denotes a chemical moiety that is capable of binding a capture reagent. Epoxide and carbodiimidazole are useful reactive moieties to covalently bind polypeptide capture reagents such as antibodies or cellular receptors. Nitriloacetic acid and iminodiacetic acid are useful reactive moieties that function as chelating agents to bind metal ions that interact non-covalently with histidine containing peptides. A "reactive surface" is a surface to which a reactive moiety is bound. An "adsorbent" or "capture reagent" can be any material capable of

binding a biomarker of the invention. Suitable adsorbents for use in SELDI, according to the invention, are described in U.S. patent No. 6,225,047, *supra*.

[0071] One type of adsorbent is a "chromatographic adsorbent," which is a material typically used in chromatography. Chromatographic adsorbents include, for example, ion exchange materials, metal chelators, immobilized metal chelates, hydrophobic interaction adsorbents, hydrophilic interaction adsorbents, dyes, simple biomolecules (e.g., nucleotides, amino acids, simple sugars and fatty acids), mixed mode adsorbents (e.g., hydrophobic attraction/electrostatic repulsion adsorbents).

"Biospecific adsorbent" is another category, for adsorbents that contain a biomolecule, e.g., a nucleotide, a nucleic acid molecule, an amino acid, a polypeptide, a polysaccharide, a lipid, a steroid or a conjugate of these (e.g., a glycoprotein, a lipoprotein, a glycolipid). In certain instances the biospecific adsorbent can be a macromolecular structure such as a multiprotein complex, a biological membrane or a virus. Illustrative biospecific adsorbents are antibodies, receptor proteins, and nucleic acids. A biospecific adsorbent typically has higher specificity for a target analyte than a chromatographic adsorbent.

[0072] Another version of SELDI is Surface-Enhanced Neat Desorption (SEND), which involves the use of probes comprising energy absorbing molecules that are chemically bound to the probe surface ("SEND probe"). The phrase "Energy absorbing molecules" (EAM) denotes molecules that are capable of absorbing energy from a laser desorption ionization source and, thereafter, contributing to desorption and ionization of analyte molecules in contact therewith. The EAM category includes molecules used in MALDI, frequently referred to as "matrix," and is exemplified by cinnamic acid derivatives, sinapinic acid (SPA), cyano-hydroxy-cinnamic acid (CHCA) and dihydroxybenzoic acid, ferulic acid, and hydroxyaceto-phenone derivatives. The category also includes EAMs used in SELDI, as enumerated, for example, by U.S. 5,719,060 and U.S. 60/351,971 (Kitagawa), filed January 25, 2002.

[0073] Another version of SELDI, called Surface-Enhanced Photolabile Attachment and Release (SEPAR), involves the use of probes having moieties attached to the surface that can covalently bind an analyte, and then release the analyte through breaking a photolabile bond in the moiety after exposure to light, e.g., to laser light.

For instance, see U.S. 5,719,060. SEPAR and other forms of SELDI are readily adapted to detecting a biomarker or biomarker profile, pursuant to the present invention.

[0074] The detection of the biomarkers according to the invention can be enhanced by using certain selectivity conditions, *e.g.*, adsorbents or washing solutions. The phrase "wash solution" refers to an agent, typically a solution, which is used to affect or modify adsorption of an analyte to an adsorbent surface and/or to remove unbound materials from the surface. The elution characteristics of a wash solution can depend, for example, on pH, ionic strength, hydrophobicity, degree of chaotropism, detergent strength, and temperature.

[0075] Pursuant to one aspect of the present invention, a sample is analyzed by means of a "biochip," a term that denotes a solid substrate, having a generally planar surface, to which a capture reagent (adsorbent) is attached. Frequently, the surface of a biochip comprises a plurality of addressable locations, each of which has the capture reagent bound there. A biochip can be adapted to engage a probe interface and, hence, function as a probe in gas phase ion spectrometry preferably mass spectrometry. Alternatively, a biochip of the invention can be mounted onto another substrate to form a probe that can be inserted into the spectrometer.

[0076] A variety of biochips is available for the capture of biomarkers, in accordance with the present invention, from commercial sources such as Ciphergen Biosystems (Fremont, CA), Perkin Elmer (Packard BioScience Company (Meriden CT), Zyomyx (Hayward, CA), and Phyllos (Lexington, MA). Exemplary of these biochips are those described in U.S. patents No. 6,225,047, *supra*, and No. 6,329,209 (Wagner *et al.*), and in PCT publications WO 99/51773 (Kuimelis and Wagner) and WO 00/56934 (Englert *et al.*).

[0077] More specifically, biochips produced by Ciphergen Biosystems have surfaces, presented on an aluminum substrate in strip form, to which are attached, at addressable locations, chromatographic or biospecific adsorbents. The surface of the strip is coated with silicon dioxide.

[0078] Illustrative of Ciphergen ProteinChip® arrays are biochips H4, SAX-2, WCX-2, and IMAC-3, which include a functionalized, cross-linked polymer in the

form of a hydrogel, physically attached to the surface of the biochip or covalently attached through a silane to the surface of the biochip. The H4 biochip has isopropyl functionalities for hydrophobic binding. The SAX-2 biochip has quaternary ammonium functionalities for anion exchange. The WCX-2 biochip has carboxylate functionalities for cation exchange. The IMAC-3 biochip has nitriloacetic acid functionalities that adsorb transition metal ions, such as Cu^{++} and Ni^{++} , by chelation. These immobilized metal ions, in turn, allow for adsorption of biomarkers by coordinate covalent bonding. Thus, CIPHERGEN's IMAC ProteinChip® arrays are sold with reactive moieties that become adsorbent upon the addition by the user of a metal solution.

[0079] In keeping with the above-described principles, a substrate with an adsorbent is contacted with the sample, containing serum, for a period of time sufficient to allow biomarker that may be present to bind to the adsorbent. In one embodiment of the invention, more than one type of substrate with adsorbent thereon is contacted with the biological sample. For example, a sample may be applied to both a WCX and an IMAC chip. This technique can allow for even more definitive assessment of cancer status. After the incubation period, the substrate is washed to remove unbound material. Any suitable washing solutions can be used; preferably, aqueous solutions are employed.

[0080] An energy absorbing molecule then is applied to the substrate with the bound biomarkers. As noted, an energy absorbing molecule is a molecule that absorbs energy from an energy source such as a laser, thereby assisting in desorption of biomarkers from the substrate. Exemplary energy absorbing molecules include, as noted above, cinnamic acid derivatives, sinapinic acid and dihydroxybenzoic acid. Preferably sinapinic acid is used.

[0081] The biomarkers bound to the substrates are detected in a gas phase ion spectrometer such as a time-of-flight mass spectrometer. The biomarkers are ionized by an ionization source such as a laser, the generated ions are collected by an ion optic assembly, and then a mass analyzer disperses and analyzes the passing ions. The detector then translates information of the detected ions into mass-to-charge

ratios. Detection of a biomarker typically will involve detection of signal intensity. Thus, both the quantity and mass of the biomarker can be determined.

[0082] Data generated by desorption and detection of biomarkers can be analyzed with the use of a programmable digital computer. The computer program analyzes the data to indicate the number of markers detected, and optionally the strength of the signal and the determined molecular mass for each biomarker detected. Data analysis can include steps of determining signal strength of a biomarker and removing data deviating from a predetermined statistical distribution. For example, the observed peaks can be normalized, by calculating the height of each peak relative to some reference. The reference can be background noise generated by the instrument and chemicals such as the energy absorbing molecule which is set as zero in the scale.

[0083] The computer can transform the resulting data into various formats for display. The standard spectrum can be displayed, but in one useful format only the peak height and mass information are retained from the spectrum view, yielding a cleaner image and enabling biomarkers with nearly identical molecular weights to be more easily seen. In another useful format, two or more spectra are compared, conveniently highlighting unique biomarkers and biomarkers that are up- or down-regulated between samples. Using any of these formats, one can readily determine whether a particular biomarker is present in a sample.

[0084] Software used to analyze the data can include code that applies an algorithm to the analysis of the signal to determine whether the signal represents a peak in a signal that corresponds to a biomarker according to the present invention. The software also can subject the data regarding observed biomarker peaks to classification tree or ANN analysis, to determine whether a biomarker peak or combination of biomarker peaks is present that indicates lung cancer status. Analysis of the data may be "keyed" to a variety of parameters that are obtained either directly or indirectly from the mass spectrometric analysis of the sample. These parameters include, but are not limited to, the presence or absence of one or more peaks, the height of one or more peaks, the log of the height of one or more peaks, and other arithmetic manipulations of peak height data.

[0085] In another aspect, the present invention provides kits for aiding in the diagnosis of lung cancer status, which kits are used to detect biomarkers according to the invention. The kits screen for the presence of biomarkers and combinations of biomarkers that are differentially present in samples from normal subjects and subjects with lung cancer.

[0086] In one embodiment, the kit comprises a substrate having an adsorbent thereon, wherein the adsorbent is suitable for binding a biomarker according to the invention, and a washing solution or instructions for making a washing solution, in which the combination of the adsorbent and the washing solution allows detection of the biomarker using gas phase ion spectrometry, e.g., mass spectrometry. The kit may include more than type of adsorbent, each present on a different substrate.

[0087] In another embodiment, a kit of the invention may include a first substrate, comprising an adsorbent thereon, and a second substrate onto which the first substrate is positioned to form a probe, which can be inserted into a gas phase ion spectrometer, e.g., a mass spectrometer. In another embodiment, an inventive kit may comprise a single substrate that can be inserted into the spectrometer.

[0088] In a further embodiment, such a kit can comprise instructions for suitable operational parameters in the form of a label or separate insert. For example, the instructions may inform a consumer how to collect the sample or how to wash the probe. In yet another embodiment the kit can comprise one or more containers with biomarker samples, to be used as standard(s) for calibration.

[0089] In a preferred embodiment, the detection of biomarkers for diagnosis of lung cancer in a subject entails contacting a sample from a subject or patient, preferably a serum sample, with a substrate having an adsorbent thereon under conditions that allow binding between the biomarker and the adsorbent, and then detecting the biomarker bound to the adsorbent by gas phase ion spectrometry, preferably by Surface Enhanced Laser Desorption/Ionization (SELDI) mass spectrometry. The biomarkers are ionized by an ionization source such as a laser. The generated ions are collected by an ion optic assembly and accelerated toward an ion detector. Ions that strike the detector generate an electric potential that is digitized by a high speed time-array recording device that digitally captures the analog signal. CIPHERGEN's

ProteinChip® system employs an analog-to-digital converter (ADC) to accomplish this. The ADC integrates detector output at regularly spaced time intervals into time-dependent bins. The time intervals typically are one to four nanoseconds long. Furthermore, the time-of-flight spectrum ultimately analyzed typically does not represent the signal from a single pulse of ionizing energy against a sample, but rather the sum of signals from a number of pulses. This reduces noise and increases dynamic range. This time-of-flight data is then subject to data processing. In CIPHERGEN's ProteinChip® software, data processing typically includes TOF-to-M/Z transformation, baseline subtraction, high frequency noise filtering. Thus, both the quantity and mass of the biomarker can be determined.

[0090] The detection of the biomarkers can be enhanced by using certain selectivity conditions, *e.g.*, adsorbents or washing solutions. In one embodiment, the same or similar selectivity conditions that were used to discover the biomarkers are used in the method of detecting the biomarker in the sample. For example, immobilized metal affinity capture chips such as the Cu(II) IMAC3 and weak cationic exchange chips such as the WCX2 chips are preferred as the adsorbents for biomarker detection. However, other adsorbents can be used, as long as they have the binding characteristics suitable for binding the biomarkers.

[0091] More particularly, armed with the information regarding the biomarkers identified herein, various methods can be used to recognize patterns of doublets, triplets, and higher combinations of biomarkers according to the invention. These methods take raw data regarding which peaks are present and their intensity and provide a differential diagnosis of lung cancer versus normal for a sample.

[0092] Thus, the process can be divided into the learning phase and the classification phase. In the learning phase, a learning algorithm is applied to a data set that includes members of the different classes that are meant to be classified, for example, data from a plurality of samples diagnosed as cancer and data from a plurality of samples assigned a negative diagnosis. The methods used to analyze the data include, but are not limited to, artificial neural network, support vector machines, genetic algorithm and self-organizing maps and classification and regression tree analysis. These methods are described, for example, in WO 01/31579, May 3, 2001

(Barnhill *et al.*); WO 02/06829, January 24, 2002 (Hitt *et al.*) and WO 02/42733, May 30, 2002 (Paulse *et al.*). The learning algorithm produces a classifying algorithm. The classifier is keyed to elements of the data, such as particular markers and particular intensities of markers, usually in combination, that can classify an unknown sample into one of the two classes. The classifier is ultimately used for diagnostic testing.

[0093] Software, both freeware and proprietary software, is readily available to analyze such patterns in data, and to devise additional patterns with any predetermined criteria for success. Those biomarkers which by themselves are predictive of a differential diagnosis of lung cancer versus normal do not require pattern recognition software to analyze the data.

[0094] The following examples are offered by way of illustration, and are not limiting.

Example I. Fractionation of serum

Buffers:

1. U9 (9M urea, 2% CHAPS, 50mM Tris-HCl pH9)
2. U1 (1M urea, 0.22% CHAPS, 50mM Tris-HCl pH9)
3. wash buffer 1: 50mM Tris-HCl with 0.1 % n-octyl β -D-Glucopyranoside (OGP) pH9
4. wash buffer 2: 100mM sodium phosphate with 0.1% OGP pH7
5. wash buffer 3: 100mM sodium acetate with 0.1% OGP pH5
6. wash buffer 4: 100mM sodium acetate with 0.1% OGP pH4
7. wash buffer 5: 50mM sodium citrate with 0.1% OGP pH3
8. wash buffer 6: 33.3% isopropanol / 16.7% acetonitrile / 0.1 % trifluoroacetic acid in water.

[0095] Thirty microliters of U9 buffer were added to 20 μ L of serum in a tube and were mixed at 4°C for 20 minutes. Ion exchange resin (Q Ceramic HyperDF ion exchange resin, Biosepra SA, France) was washed 3 times with 5 bed volumes of 50mM Tris-HCl pH9 and stored in 50% suspension. To each well of a 96-well filter plate (96-well Silent Screen filter plate, Lonrodvne membrane, 0.45 micron pore,

Nalge Nunc International, USA), 125 μ L of ion exchange resin (50% suspension) was added on a Biomek 2000 Automation Workstation (Beckman Coulter, Fullerton, CA), washed 3 times with 150 μ L U1 buffer, and vacuum dried. Urea-treated serum was transferred to each well of ion exchange resin. The serum tube was rinsed with 50 μ L of U1 buffer, which was also transferred to the corresponding well in filter plate. The filter plate was mixed on a platform shaker at 4°C for 30 minutes. Flow-through fraction was collected in a 96-well plate by vacuum suction (Fraction 1). Then, 100 μ L of wash buffer 1 was added to each well of filter plate and mixed for 10 minutes at room temperature. Eluant was collected into the same 96-well plate (Fraction 1). Resins in the filter plate were subsequently washed two times each with 100 μ L wash buffers 2, 3, 4, 5 and 6. Each eluant (total volume of 200 μ L) was collected in a 96-well plate (Fractions 2,3,4,5 and 6).

Example 2. SELDI analysis of fractionated serum

[0096] ProteinChip® Arrays were set up in 96-well bioprocessors. Buffer delivery and sample incubation were performed on a Biomek 2000 Automation Workstation. Each serum fraction was analyzed on IMAC3 (loaded with copper) and WCX2 ProteinChip® Arrays in duplicates. IMAC3 copper and WCX2 arrays (CIPHERGEN Biosystems Inc, Fremont, CA) were equilibrated two times with 150 μ L of binding buffer (100mM sodium phosphate + 0.5M NaCl pH7 for IMAC3, 100mM sodium acetate pH4 for WCX2). Each serum fraction was diluted in the corresponding binding buffer (1/5 dilution for IMAC3 and 1/10 dilution for WCX2) and 100 μ L was applied to each ProteinChip® array. Incubation was performed on a platform shaker at room temperature for 30 minutes. Each array was washed three times with 150 μ L of corresponding binding buffer and rinsed two times with water. ProteinChip® arrays were air-dried. Sinapinic acid matrix (prepared in 50% acetonitrile, 0.5% trifluoroacetic acid) was applied to each array. ProteinChip® arrays were read on a ProteinChip® PBSII Reader (CIPHERGEN Biosystems Inc.) A total of 253 laser shots were averaged for each array.

[0097] All publications and patent documents cited in this application are incorporated by reference in their entirety for all purposes to the same extent as if

each individual publication or patent document were so individually denoted. By their citation of various references in this document Applicants do not admit that any particular reference is "prior art" to their invention.

What we claim is:

1. A method for qualifying lung carcinoma status in a subject, comprised of analyzing a biological sample from said subject for a diagnostic level of a protein selected from either a first group consisting of

(i) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, IM-155, IM-157, IM-176, IM-445, IM-177, IM-440, IM-468, IM-438, IM-547, IM-359, IM-436, IM-106, IM-455, IM-444, IM-158, IM-265, IM-50, IM-159, IM-156, IM-439, IM-157, IM-508, IM-514, IM-478, IM-473, IM-360, IM-435, IM-150, IM-151, IM-110, IM-51, IM-163, IM-437, IM-546, IM-153, and IM-268,

or from a second group consisting of

(ii) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, WM-70, WM-369, WM-17, WM-473, WM-47, WM-203, WM-276, WM-279, WM-62, WM-366, WM-456, WM-428, WM-384, WM-287, WM-420, WM-292, WM-431, WM-455, WM-20, WM-340, WM-105, WM-389, WM-63, WM-354, WM-450, WM-466, WM-296, WM-343, WM-341, WM-339, WM-55, WM-66, WM-48, WM-38, WM-138, and WM-310;

wherein the biomarker is differentially present in samples of a subject with lung cancer and a normal subject that is free of lung cancer.

2. The method according to claim 1, wherein the protein is selected from either a first group consisting of

(i) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, and IM-155,

or from a second group consisting of

(ii) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, and WM-70.

3. The method according to claim 1, wherein the protein is selected from either a first group consisting of

(i) IM-522, IM-273, IM-520, IM-519, and IM-454,

or from a second group consisting

(ii) WM-61, WM-447, WM-446, WM-133, and WM-119.

4. The method according to claim 1, which uses a single biomarker selected from the group consisting of the WM-446 and WM-447.

5. A method for qualifying lung carcinoma risk in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from either a first group consisting of

(i) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, IM-155, IM-157, IM-176, IM-445, IM-177, IM-440, IM-468, IM-438, IM-547, IM-359, IM-436, IM-106, IM-455, IM-444, IM-158, IM-265, IM-50, IM-159, IM-156, IM-439, IM-157, IM-508, IM-514, IM-478, IM-473, IM-360, IM-435, IM-150, IM-151, IM-110, IM-51, IM-163, IM-437, IM-546, IM-153, and IM-268,

or from a second group consisting of

(ii) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, WM-70, WM-369, WM-17, WM-473, WM-47, WM-203, WM-276, WM-279, WM-62, WM-366, WM-456, WM-428, WM-384, WM-287, WM-420, WM-292, WM-431, WM-455, WM-20, WM-340, WM-105, WM-389, WM-63, WM-354, WM-450, WM-466, WM-296, WM-343, WM-341, WM-339, WM-55, WM-66, WM-48, WM-38, WM-138, and WM-310.

6. The method according to claim 5, wherein the pattern-recognition analysis is keyed to a pair of peaks selected either from a first group consisting of

(i) IM-266 and IM-474, IM-266 and IM-38, IM-266 and IM-454, IM-266 and IM-522, IM-266 and IM-544, IM-266 and IM-471, IM-474 and IM-151, IM-474 and IM-156, IM-474 and IM-544, IM-474 and IM-38, IM-522 and IM-507, IM-522 and IM-156, and IM-522 and IM-440;

or from a second group consisting of

(ii) WM-447 and WM-59, WM-447 and WM-19, WM-447 and WM-118, WM-447 and WM-473, WM-19 and WM-59, WM-19 and WM-473, WM-19 and WM-369, WM-61 and WM-154, WM-61 and WM-369, WM-118 and WM-59 and WM-282 and WM-127.

7. The method according to claim 5, wherein the pattern-recognition analysis is keyed to a pair of peaks selected from either a first group consisting of

(i) IM-266 and IM-474, IM-266 and IM-544, and IM-156 and IM-522;

or from a second group consisting of

(ii) WM-447 and WM-59, WM-447 and WM-19, and WM-19 and WM-59.

8. The method according to claim 5, wherein the pattern-recognition analysis is keyed to a triplet of peaks selected from

(i) IM-266, IM-454 and IM-474; and IM-266, IM-474 and IM-544;

or wherein the analysis is keyed to

(ii) WM-447, WM-19 and WM-473.

9. A kit for detecting and diagnosing lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers selected from either a first group consisting of

(i) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, IM-155, IM-157, IM-176, IM-445, IM-177, IM-440, IM-468, IM-438, IM-547, IM-359, IM-436, IM-106, IM-455, IM-444, IM-158, IM-265, IM-50, IM-159, IM-156, IM-439, IM-157, IM-508, IM-514, IM-478, IM-473, IM-360, IM-435, IM-150, IM-151, IM-110, IM-51, IM-163, IM-437, IM-546, IM-153, and IM-268,

or from a second group consisting of

(ii) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, WM-70, WM-369, WM-17, WM-473, WM-47, WM-203, WM-276, WM-279, WM-62, WM-366, WM-456, WM-428, WM-384, WM-287, WM-420, WM-292, WM-431, WM-455, WM-20, WM-340, WM-105, WM-389, WM-63, WM-354, WM-450, WM-466, WM-296, WM-343, WM-341, WM-339, WM-55, WM-66, WM-48, WM-38, WM-138, and WM-310, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

10. A kit according to claim 9, further comprising a washing solution or instructions for making a washing solution.

11. A kit according to claim 9, wherein the substrate is a SELDI probe that comprises either (i) functionalities that adsorb transition metal ions by chelation or (ii) functionalities that allow for cation exchange.

12. A method for qualifying lung adenocarcinoma status in a subject, comprised of analyzing a biological sample from said subject for a level of a protein selected from the group consisting of WM-447, WM-652, WM-61, WM-446, WM-290, WM-363, WM-133, WM-341, WM-285, WM-366, WM-282, WM-362, WM-310, WM-292, WM-120, WM-134, WM-276, WM-428, WM-277, WM-20, WM-119, WM-340, WM-48, WM-389, WM-450, WM-47, WM-343, WM-17, WM-583, WM-70, WM-706, WM-346, WM-466, WM-646, WM-384, WM-336, WM-294, WM-339, WM-473, WM-369, WM-38, WM-283, WM-685, WM-66, WM-55, WM-650, WM-307, WM-278, WM-342, and WM-429.

13. The method according to claim 12, wherein the protein is selected from the group consisting of WM-447, WM-652, WM-61, WM-446, WM-290, WM-363, WM-133, WM-341, WM-285, WM-366, WM-282, WM-362, WM-310, WM-292, and WM-120.

14. The method according to claim 12, wherein the protein is selected from the group consisting of WM-447, WM-652, WM-61, WM-446, WM-290.

15. A method for qualifying status of lung adenocarcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from either a first group consisting of WM-447, WM-652, WM-61, WM-446, WM-290, WM-363, WM-133, WM-341, WM-285, WM-366, WM-282, WM-362, WM-310, WM-292, WM-120, WM-134, WM-276, WM-428, WM-277, WM-20, WM-119, WM-340, WM-48, WM-

389, WM-450, WM-47, WM-343, WM-17, WM-583, WM-70, WM-706, WM-346, WM-466, WM-646, WM-384, WM-336, WM-294, WM-339, WM-473, WM-369, WM-38, WM-283, WM-685, WM-66, WM-55, WM-650, WM-307, WM-278, WM-342, and WM-429.

16. The method according to claim 15, wherein the protein is selected from the group consisting of WM-447, WM-652, WM-61, WM-446, WM-290, WM-363, WM-133, WM-341, WM-285, WM-366, WM-282, WM-362, WM-310, WM-292, and WM-120.

17. The method according to claim 15, wherein the protein is selected from the group consisting of WM-447, WM-652, WM-61, WM-446, WM-290.

18. A kit for detecting and diagnosing lung adenocarcinoma, comprising
(A) an adsorbent attached to a substrate that retains one or more of biomarkers selected from the group consisting of WM-447, WM-652, WM-61, WM-446, WM-290, WM-363, WM-133, WM-341, WM-285, WM-366, WM-282, WM-362, WM-310, WM-292, WM-120, WM-134, WM-276, WM-428, WM-277, WM-20, WM-119, WM-340, WM-48, WM-389, WM-450, WM-47, WM-343, WM-17, WM-583, WM-70, WM-706, WM-346, WM-466, WM-646, WM-384, WM-336, WM-294, WM-339, WM-473, WM-369, WM-38, WM-283, WM-685, WM-66, WM-55, WM-650, WM-307, WM-278, WM-342, and WM-429, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

19. A kit according to claim 18, further comprising a washing solution or instructions for making a washing solution.

20. A kit according to claim 18, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

21. A method for qualifying squamous cell lung carcinoma status in a subject, comprised of analyzing a biological sample from said subject for a level of a protein selected from the group consisting of WM-447, WM-61, WM-277, WM-446, WM-133, WM-134, WM-363, WM-362, WM-276, WM-706, WM-203, WM-466, WM-366, WM-65, WM-70, WM-341, WM-429, WM-347, WM-17, WM-47, WM-431, WM-62, WM-473, WM-384, WM-438, WM-652, WM-282, WM-389, WM-290,

WM-278, WM-456, WM-673, WM-340, WM-55, WM-455, WM-645, WM-138, WM-420, WM-450, WM-369, WM-279, WM-342, WM-471, WM-674, WM-120, WM-20, WM-287, WM-83, WM-154, and WM-128.

22. The method according to claim 21, wherein the protein is selected from the group consisting of WM-447, WM-61, WM-277, WM-446, WM-133, WM-134, WM-363, WM-362, WM-276, WM-706, WM-203, WM-466, WM-366, WM-65, and WM-70.

23. The method according to claim 21, wherein the protein is selected from the group consisting of WM-447, WM-61, WM-277, WM-446, and WM-133.

24. A method for qualifying status of squamous cell lung carcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from either a first group consisting of WM-447, WM-61, WM-277, WM-446, WM-133, WM-134, WM-363, WM-362, WM-276, WM-706, WM-203, WM-466, WM-366, WM-65, WM-70, WM-341, WM-429, WM-347, WM-17, WM-47, WM-431, WM-62, WM-473, WM-384, WM-438, WM-652, WM-282, WM-389, WM-290, WM-278, WM-456, WM-673, WM-340, WM-55, WM-455, WM-645, WM-138, WM-420, WM-450, WM-369, WM-279, WM-342, WM-471, WM-674, WM-120, WM-20, WM-287, WM-83, WM-154, and WM-128.

25. The method according to claim 24, wherein the protein is selected from the group consisting of WM-447, WM-61, WM-277, WM-446, WM-133, WM-134, WM-363, WM-362, WM-276, WM-706, WM-203, WM-466, WM-366, WM-65, and WM-70.

26. The method according to claim 24, wherein the protein is selected from the group consisting of WM-447, WM-61, WM-277, WM-446, and WM-133.

27. A kit for detecting and diagnosing squamous cell lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers selected from the group consisting of WM-447, WM-61, WM-277, WM-446, WM-133, WM-134, WM-363, WM-362, WM-276, WM-706, WM-203, WM-466, WM-366, WM-65, WM-70, WM-341, WM-429, WM-347, WM-17, WM-47, WM-431, WM-62, WM-473, WM-384, WM-438, WM-652, WM-282, WM-389, WM-290, WM-278, WM-456, WM-673, WM-340, WM-55, WM-455, WM-645, WM-138, WM-420, WM-450, WM-369, WM-279, WM-342, WM-471, WM-674, WM-120, WM-20, WM-287, WM-83, WM-154, and WM-128, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

28. A kit according to claim 27, further comprising a washing solution or instructions for making a washing solution.

29. A kit according to claim 27, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

30. A method for qualifying small cell lung carcinoma status in a subject, comprised of analyzing a biological sample from said subject for a level of a protein selected from the group consisting of WM-70, WM-706, WM-369, WM-447, WM-61, WM-652, WM-282, WM-446, WM-456, WM-134, WM-203, WM-646, WM-455, WM-65, WM-685, WM-473, WM-343, WM-466, WM-341, WM-340, WM-363, WM-339, WM-457, WM-86, WM-506, WM-72, WM-287, WM-82, WM-528, WM-85, WM-73, WM-138, WM-384, WM-83, WM-450, WM-310, WM-277, WM-79, WM-207, WM-278, WM-290, WM-366, WM-472, WM-420, WM-147, WM-55, WM-669, WM-357, WM-429, and WM-279.

31. The method according to claim 30, wherein the protein is selected from the group consisting of WM-70, WM-706, WM-369, WM-447, WM-61, WM-652, WM-282, WM-446, WM-456, WM-134, WM-203, WM-646, WM-455, WM-65, and WM-685.

32. The method according to claim 30, wherein the protein is selected from the group consisting of WM-70, WM-706, WM-369, WM-447, and WM-61.

33. A method for qualifying status of small cell lung carcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from either a first group consisting of WM-70, WM-706, WM-369, WM-447, WM-61, WM-652, WM-282, WM-446, WM-456, WM-134, WM-203, WM-646, WM-455, WM-65, WM-685, WM-473, WM-343, WM-466, WM-341, WM-340, WM-363, WM-339, WM-457, WM-86, WM-506, WM-72, WM-287, WM-82, WM-528, WM-85, WM-73, WM-138, WM-384, WM-83, WM-450, WM-310, WM-277, WM-79, WM-207, WM-278, WM-290, WM-366, WM-472, WM-420, WM-147, WM-55, WM-669, WM-357, WM-429, and WM-279.

34. The method according to claim 33, wherein the protein is selected from the group consisting of WM-70, WM-706, WM-369, WM-447, WM-61, WM-652, WM-282, WM-446, WM-456, WM-134, WM-203, WM-646, WM-455, WM-65, and WM-685.

35. The method according to claim 33, wherein the protein is selected from the group consisting of WM-70, WM-706, WM-369, WM-447, and WM-61.

36. A kit for detecting and diagnosing small cell lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers selected from the group consisting of WM-70, WM-706, WM-369, WM-447, WM-61, WM-652, WM-282, WM-446, WM-456, WM-134, WM-203, WM-646, WM-455, WM-65, WM-685, WM-473, WM-343, WM-466, WM-341, WM-340, WM-363, WM-339, WM-457, WM-86, WM-506, WM-72, WM-287, WM-82, WM-528, WM-85, WM-73, WM-138, WM-384, WM-83, WM-450, WM-310, WM-277, WM-79, WM-207, WM-278, WM-290, WM-366, WM-472, WM-420, WM-147, WM-55, WM-669, WM-357, WM-429, and WM-279, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

37. A kit according to claim 36, further comprising a washing solution or instructions for making a washing solution.

38. A kit according to claim 36, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

39. A method for qualifying non-small cell lung carcinoma status in a subject, comprised of analyzing a biological sample from said subject for a level of a protein selected from the group consisting of WM-341, WM-342, WM-343, WM-48, WM-340, WM-346, WM-47, WM-339, WM-389, WM-669, WM-447, WM-652, WM-154, WM-587, WM-456, WM-450, WM-283, WM-207, WM-436, WM-384, WM-61, WM-167, WM-382, WM-285, WM-650, WM-203, WM-119, WM-282, WM-686, WM-383, WM-429, WM-11, WM-208, WM-451, WM-473, WM-220, WM-685, WM-338, WM-71, WM-266, WM-70, WM-545, WM-675, WM-446, WM-120, WM-267, WM-466, WM-347, WM-153, and WM-38.

40. The method according to claim 39, wherein the protein is selected from the group consisting of WM-341, WM-342, WM-343, WM-48, WM-340, WM-346, WM-47, WM-339, WM-389, WM-669, WM-447, WM-652, WM-154, WM-587, and WM-456.

41. The method according to claim 39, wherein the protein is selected from the group consisting of WM-341, WM-342, WM-343, WM-48, and WM-340.

42. A method for qualifying status of non-small cell lung carcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from the group consisting of WM-341, WM-342, WM-343, WM-48, WM-340, WM-346, WM-47, WM-339, WM-389, WM-669, WM-447, WM-652, WM-154, WM-587, WM-456, WM-450, WM-283, WM-207, WM-436, WM-384, WM-61, WM-167, WM-382, WM-285, WM-650, WM-203, WM-119, WM-282, WM-686, WM-383, WM-429, WM-11, WM-208, WM-451, WM-473, WM-220, WM-685, WM-338, WM-71, WM-266, WM-70, WM-545, WM-675, WM-446, WM-120, WM-267, WM-466, WM-347, WM-153, and WM-38.

43. The method according to claim 42, wherein the protein is selected from the group consisting of WM-341, WM-342, WM-343, WM-48, WM-340, WM-346, WM-47, WM-339, WM-389, WM-669, WM-447, WM-652, WM-154, WM-587, and WM-456.

44. The method according to claim 42, wherein the protein is selected from the group consisting of WM-341, WM-342, WM-343, WM-48, and WM-340.

45. A kit for detecting and diagnosing non-small cell lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers WM-341, WM-342, WM-343, WM-48, WM-340, WM-346, WM-47, WM-339, WM-389, WM-669, WM-447, WM-652, WM-154, WM-587, WM-456, WM-450, WM-283, WM-207, WM-436, WM-384, WM-61, WM-167, WM-382, WM-285, WM-650, WM-203, WM-119, WM-282, WM-686, WM-383, WM-429, WM-11, WM-208, WM-451, WM-473, WM-220, WM-685, WM-338, WM-71, WM-266, WM-70, WM-545, WM-675, WM-446, WM-120, WM-267, WM-466, WM-347, WM-153, and WM-38, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

46. A kit according to claim 45, further comprising a washing solution or instructions for making a washing solution.

47. A kit according to claim 45, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

48. A method for qualifying large cell lung carcinoma status in a subject, comprised of analyzing a biological sample from said subject for a level of a protein selected from the group consisting of WM-16, WM-26, WM-499, WM-134, WM-647, WM-277, WM-310, WM-363, WM-446, WM-221, WM-648, WM-657, WM-290, WM-328, WM-447, WM-684, WM-183, WM-190, WM-686, WM-397, WM-466, WM-20, WM-17, WM-545, WM-47, WM-191, WM-147, WM-480, WM-590, WM-218, WM-285, WM-652, WM-651, WM-366, WM-403, WM-418, WM-430, WM-456, WM-714, WM-646, WM-109, WM-302, WM-587, WM-375, WM-131, WM-706, WM-398, WM-309, WM-55, and WM-488.

49. The method according to claim 48, wherein the protein is selected from the group consisting of WM-16, WM-26, WM-499, WM-134, WM-647, WM-277, WM-310, WM-363, WM-446, WM-221, WM-648, WM-657, WM-290, WM-328, and WM-447.

50. The method according to claim 48, wherein the protein is selected from the group consisting of WM-16, WM-26, WM-499, WM-134, and WM-647.

51. A method for qualifying status of large cell lung carcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from the group consisting of WM-16, WM-26, WM-499, WM-134, WM-647, WM-277, WM-310, WM-363, WM-446, WM-221, WM-648, WM-657, WM-290, WM-328, WM-447, WM-684, WM-183, WM-190, WM-686, WM-397, WM-466, WM-20, WM-17, WM-545, WM-47, WM-191, WM-147, WM-480, WM-590, WM-218, WM-285, WM-652, WM-651, WM-366, WM-403, WM-418, WM-430, WM-456, WM-714, WM-646, WM-109, WM-302, WM-587, WM-375, WM-131, WM-706, WM-398, WM-309, WM-55, and WM-488.

52. The method according to claim 51, wherein the protein is selected from the group consisting of WM-16, WM-26, WM-499, WM-134, WM-647, WM-277, WM-310, WM-363, WM-446, WM-221, WM-648, WM-657, WM-290, WM-328, and WM-447.

53. The method according to claim 51, wherein the protein is selected from the group consisting of WM-16, WM-26, WM-499, WM-134, and WM-647.

54. A kit for detecting and diagnosing large cell lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers WM-16, WM-26, WM-499, WM-134, WM-647, WM-277, WM-310, WM-363, WM-446, WM-221, WM-648, WM-657, WM-290, WM-328, WM-447, WM-684, WM-183, WM-190, WM-686, WM-397, WM-466, WM-20, WM-17, WM-

545, WM-47, WM-191, WM-147, WM-480, WM-590, WM-218, WM-285, WM-652, WM-651, WM-366, WM-403, WM-418, WM-430, WM-456, WM-714, WM-646, WM-109, WM-302, WM-587, WM-375, WM-131, WM-706, WM-398, WM-309, WM-55, and WM-488, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

55. A kit according to claim 50, further comprising a washing solution or instructions for making a washing solution.

56. A kit according to claim 50, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

57. A method for distinguishing lung adenocarcinoma from squamous lung carcinoma in a subject, comprised of analyzing a biological sample from said subject for a level of a protein selected from the group consisting of WM-62, WM-415, WM-152, WM-385, WM-347, WM-134, WM-36, WM-108, WM-99, WM-151, WM-289, WM-363, WM-61, WM-117, WM-211, WM-362, WM-133, WM-414, WM-277, WM-141, WM-64, WM-135, WM-447, WM-383, WM-338, WM-63, WM-142, WM-446, WM-186, WM-111, WM-445, WM-455, WM-276, WM-444, WM-181, WM-35, WM-285, WM-456, WM-39, WM-82, WM-17, WM-203, WM-83, WM-412, WM-96, WM-74, WM-457, WM-431, WM-340, and WM-49.

58. The method according to claim 57, wherein the protein is selected from the group consisting of WM-62, WM-415, WM-152, WM-385, WM-347, WM-134, WM-36, WM-108, WM-99, WM-151, WM-289, WM-363, WM-61, WM-117, and WM-211.

59. The method according to claim 57, wherein the protein is selected from the group consisting of WM-62, WM-415, WM-152, WM-385, and WM-347.

60. A method for distinguishing lung adenocarcinoma from squamous lung carcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from the group

consisting of WM-62, WM-415, WM-152, WM-385, WM-347, WM-134, WM-36, WM-108, WM-99, WM-151, WM-289, WM-363, WM-61, WM-117, WM-211, WM-362, WM-133, WM-414, WM-277, WM-141, WM-64, WM-135, WM-447, WM-383, WM-338, WM-63, WM-142, WM-446, WM-186, WM-111, WM-445, WM-455, WM-276, WM-444, WM-181, WM-35, WM-285, WM-456, WM-39, WM-82, WM-17, WM-203, WM-83, WM-412, WM-96, WM-74, WM-457, WM-431, WM-340, and WM-49.

61. The method according to claim 60, wherein the protein is selected from the group consisting of WM-62, WM-415, WM-152, WM-385, WM-347, WM-134, WM-36, WM-108, WM-99, WM-151, WM-289, WM-363, WM-61, WM-117, and WM-211.

62. The method according to claim 60, wherein the protein is selected from the group consisting of WM-62, WM-415, WM-152, WM-385, and WM-347.

63. A kit for distinguishing lung adenocarcinoma from squamous lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers WM-16, WM-26, WM-499, WM-134, WM-647, WM-277, WM-310, WM-363, WM-446, WM-221, WM-648, WM-657, WM-290, WM-328, WM-447, WM-684, WM-183, WM-190, WM-686, WM-397, WM-466, WM-20, WM-17, WM-545, WM-47, WM-191, WM-147, WM-480, WM-590, WM-218, WM-285, WM-652, WM-651, WM-366, WM-403, WM-418, WM-430, WM-456, WM-714, WM-646, WM-109, WM-302, WM-587, WM-375, WM-131, WM-706, WM-398, WM-309, WM-55, and WM-488, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

64. A kit according to claim 63, further comprising a washing solution or instructions for making a washing solution.

65. A kit according to claim 63, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

66. A method for distinguishing lung adenocarcinoma from small cell lung carcinoma in a subject, comprised of analyzing a biological sample from said subject

for a level of a protein selected from the group consisting of WM-457, WM-72, WM-369, WM-78, WM-79, WM-73, WM-64, WM-320, WM-419, WM-85, WM-82, WM-53, WM-412, WM-440, WM-455, WM-313, WM-456, WM-86, WM-70, WM-246, WM-360, WM-190, WM-418, WM-83, WM-257, WM-138, WM-47, WM-252, WM-282, WM-60, WM-68, WM-325, WM-402, WM-411, WM-405, WM-75, WM-417, WM-387, WM-26, WM-410, WM-420, WM-164, WM-67, WM-66, WM-391, WM-340, WM-428, WM-198, WM-312, and WM-152.

67. The method according to claim 66, wherein the protein is selected from the group consisting of WM-457, WM-72, WM-369, WM-78, WM-79, WM-73, WM-64, WM-320, WM-419, WM-85, WM-82, WM-53, WM-412, WM-440, and WM-455.

68. The method according to claim 66, wherein the protein is selected from the group consisting of WM-457, WM-72, WM-369, WM-78, and WM-79.

69. A method for distinguishing lung adenocarcinoma from small cell lung carcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from either a first group consisting of WM-457, WM-72, WM-369, WM-78, WM-79, WM-73, WM-64, WM-320, WM-419, WM-85, WM-82, WM-53, WM-412, WM-440, WM-455, WM-313, WM-456, WM-86, WM-70, WM-246, WM-360, WM-190, WM-418, WM-83, WM-257, WM-138, WM-47, WM-252, WM-282, WM-60, WM-68, WM-325, WM-402, WM-411, WM-405, WM-75, WM-417, WM-387, WM-26, WM-410, WM-420, WM-164, WM-67, WM-66, WM-391, WM-340, WM-428, WM-198, WM-312, and WM-152.

70. The method according to claim 69, wherein the protein is selected from the group consisting of WM-457, WM-72, WM-369, WM-78, WM-79, WM-73, WM-64, WM-320, WM-419, WM-85, WM-82, WM-53, WM-412, WM-440, and WM-455.

71. The method according to claim 69, wherein the protein is selected from the group consisting of WM-457, WM-72, WM-369, WM-78, WM-79.

72. A kit for distinguishing lung adenocarcinoma from small cell lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers WM-276, WM-277, WM-362, WM-257, WM-363, WM-347, WM-53, WM-254, WM-17, WM-252, WM-431, WM-513, WM-446, WM-355, WM-447, WM-133, WM-245, WM-52, WM-96, WM-238, WM-243, WM-138, WM-62, WM-580, WM-134, WM-240, WM-256, WM-203, WM-111, WM-95, WM-247, WM-157, WM-242, WM-556, WM-63, WM-239, WM-234, WM-274, WM-370, WM-301, WM-449, WM-74, WM-261, WM-467, WM-237, WM-262, WM-295, WM-288, WM-384, and WM-37, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

73. A kit according to claim 72, further comprising a washing solution or instructions for making a washing solution.

74. A kit according to claim 72, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

75. A method for distinguishing squamous cell lung carcinoma from small cell lung carcinoma in a subject, comprised of analyzing a biological sample from said subject for a level of a protein selected from the group consisting of WM-276, WM-277, WM-362, WM-257, WM-363, WM-347, WM-53, WM-254, WM-17, WM-252, WM-431, WM-513, WM-446, WM-355, WM-447, WM-133, WM-245, WM-52, WM-96, WM-238, WM-243, WM-138, WM-62, WM-580, WM-134, WM-240, WM-256, WM-203, WM-111, WM-95, WM-247, WM-157, WM-242, WM-556, WM-63, WM-239, WM-234, WM-274, WM-370, WM-301, WM-449, WM-74, WM-261, WM-467, WM-237, WM-262, WM-295, WM-288, WM-384, and WM-37.

76. The method according to claim 75, wherein the protein is selected from the group consisting of WM-276, WM-277, WM-362, WM-257, WM-363, WM-347, WM-53, WM-254, WM-17, WM-252, WM-431, WM-513, WM-446, WM-355, and WM-447.

77. The method according to claim 75, wherein the protein is selected from the group consisting of WM-276, WM-277, WM-362, WM-257, and WM-363.

78. A method for distinguishing squamous cell lung carcinoma from small cell lung carcinoma in a subject, comprising

(A) providing a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, and

(B) extracting data from the spectrum and subjecting the data to pattern-recognition analysis that is keyed to at least one peak selected from either a first group consisting of WM-276, WM-277, WM-362, WM-257, WM-363, WM-347, WM-53, WM-254, WM-17, WM-252, WM-431, WM-513, WM-446, WM-355, WM-447, WM-133, WM-245, WM-52, WM-96, WM-238, WM-243, WM-138, WM-62, WM-580, WM-134, WM-240, WM-256, WM-203, WM-111, WM-95, WM-247, WM-157, WM-242, WM-556, WM-63, WM-239, WM-234, WM-274, WM-370, WM-301, WM-449, WM-74, WM-261, WM-467, WM-237, WM-262, WM-295, WM-288, WM-384, and WM-37.

79. The method according to claim 78, wherein the protein is selected from the group consisting of WM-276, WM-277, WM-362, WM-257, WM-363, WM-347, WM-53, WM-254, WM-17, WM-252, WM-431, WM-513, WM-446, WM-355, and WM-447.

80. The method according to claim 78, wherein the protein is selected from the group consisting of WM-276, WM-277, WM-362, WM-257, and WM-363.

81. A kit for distinguishing squamous cell lung carcinoma from small cell lung carcinoma, comprising

(A) an adsorbent attached to a substrate that retains one or more of the biomarkers WM-276, WM-277, WM-362, WM-257, WM-363, WM-347, WM-53, WM-254, WM-17, WM-252, WM-431, WM-513, WM-446, WM-355, WM-447, WM-133, WM-245, WM-52, WM-96, WM-238, WM-243, WM-138, WM-62, WM-580, WM-134, WM-240, WM-256, WM-203, WM-111, WM-95, WM-247, WM-157, WM-242, WM-556, WM-63, WM-239, WM-234, WM-274, WM-370, WM-301, WM-449, WM-74, WM-261, WM-467, WM-237, WM-262, WM-295, WM-288, WM-384, and WM-37, and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

82. A kit according to claim 81, further comprising a washing solution or instructions for making a washing solution.

83. A kit according to claim 81, wherein the substrate is a SELDI probe that comprises functionalities that allow for cation exchange.

84. Software for qualifying lung carcinoma status in a subject, comprising an algorithm for analyzing data extracted from a spectrum generated by mass spectroscopic analysis of a biological sample taken from the subject, wherein said data relates to one or more biomarkers selected from either a first group consisting of

(i) IM-522, IM-273, IM-520, IM-519, IM-454, IM-507, IM-521, IM-148, IM-266, IM-537, IM-471, IM-510, IM-544, IM-474, IM-155, IM-157, IM-176, IM-445, IM-177, IM-440, IM-468, IM-438, IM-547, IM-359, IM-436, IM-106, IM-455, IM-444, IM-158, IM-265, IM-50, IM-159, IM-156, IM-439, IM-157, IM-508, IM-514, IM-478, IM-473, IM-360, IM-435, IM-150, IM-151, IM-110, IM-51, IM-163, IM-437, IM-546, IM-153, and IM-268,

or from a second group consisting of

(ii) WM-61, WM-447, WM-446, WM-133, WM-119, WM-278, WM-134, WM-363, WM-282, WM-362, WM-120, WM-290, WM-65, WM-277, WM-70, WM-369, WM-17, WM-473, WM-47, WM-203, WM-276, WM-279, WM-62, WM-366, WM-456, WM-428, WM-384, WM-287, WM-420, WM-292, WM-431, WM-455, WM-20, WM-340, WM-105, WM-389, WM-63, WM-354, WM-450, WM-466, WM-296, WM-343, WM-341, WM-339, WM-55, WM-66, WM-48, WM-38, WM-138, and WM-310.

85. Software according to claim 84, wherein said algorithm carries out a pattern-recognition analysis that is keyed to data relating to at least one of the biomarkers.

86. Software according to claim 85, wherein said algorithm comprises classification tree analysis that is keyed to data relating to at least one of the biomarkers.

87. Software according to claim 85, wherein said algorithm comprises artificial neural network analysis that is keyed to data relating to at least one of the biomarkers.

88. A method for qualifying lung carcinoma status in a subject, comprised of analyzing a biological sample from said subject for a diagnostic level of a biomarker that is serum amyloid A protein or a fragment thereof.

89. A method according to claim 88, wherein said serum biomarker has an apparent molecular weight of about 2803, 3168, 3277, 3552, 3897, 4300, 4490, 4655, 5927, 6874, 7776, 7941, 8152, 8952, 9233, 10300, 10866, or 10851 Daltons.

90. A method according to claim 89, wherein said serum biomarker has an apparent molecular weight of about 3168, 3277, 3552, 3897, 4300, 4490, 4655, 7776, 7941, 8152, 8952, or 10851 Daltons.

91. A method according to claim 88, wherein said serum biomarker has an apparent molecular weight of about 11.5 to 11.7 kD.

92. A method according to claim 88, for qualifying risk of lung adenocarcinoma.

93. A method according to claim 88, for qualifying risk of squamous cell lung carcinoma.

94. A method according to claim 88, for qualifying risk of small cell lung carcinoma.

95. A method according to claim 88, for qualifying risk of non-small cell lung carcinoma.

96. A method according to claim 88, for qualifying risk of large cell lung carcinoma.

97. A kit for detecting and diagnosing lung carcinoma, comprising
(A) an adsorbent attached to a substrate that retains one or more of the biomarkers that are serum amyloid A protein or a fragment thereof.

and

(B) instructions to detect the biomarker(s) by contacting a sample with the adsorbent and detecting the biomarker(s) retained by the adsorbent.

98. A kit according to claim 97, wherein said serum biomarker has an apparent molecular weight of about 2803, 3168, 3277, 3552, 3897, 4300, 4490, 4655, 5927, 6874, 7776, 7941, 8152, 8952, 9233, 10300, 10866, or 10851 Daltons.

99. A kit according to claim 98, wherein said serum biomarker has an apparent molecular weight of about 3168, 3277, 3552, 3897, 4300, 4490, 4655, 7776, 7941, 8152, 8952, or 10851 Daltons.

100. A kit according to claim 97, wherein said serum biomarker has an apparent molecular weight of about 11.5 to 11.7 kD.

101. A kit according to claim 97, further comprising a washing solution or instructions for making a washing solution.

102. A kit according to claim 97, wherein the substrate is a SELDI probe.

FIGURE 1A

MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION
IM-1	2011	A	IM-37	3893	A	IM-72	54026	A	IM-109	2882	B	IM-110	2967	B
IM-2	2030	A	IM-38	3960	A	IM-73	60170	A	IM-111	2977	B	IM-112	2994	B
IM-3	2069	A	IM-39	3972	A	IM-75	74372	A	IM-113	3031	B	IM-114	3048	B
IM-4	2128	A	IM-40	3984	A	IM-76	75545	A	IM-115	3148	B	IM-116	3166	B
IM-5	2146	A	IM-41	4066	A	IM-77	77543	A	IM-117	3283	B	IM-118	3308	B
IM-6	2186	A	IM-42	4178	A	IM-78	79507	A	IM-119	3332	B	IM-120	3432	B
IM-7	2232	A	IM-43	4287	A	IM-79	89854	A	IM-121	3450	B	IM-122	3561	B
IM-8	2277	A	IM-44	4297	A	IM-80	101831	A	IM-123	3615	B	IM-124	3714	B
IM-9	2295	A	IM-45	4309	A	IM-81	104301	A	IM-125	3730	B	IM-126	3834	B
IM-10	2318	A	IM-46	4484	A	IM-82	125160	A	IM-127	3899	B	IM-128	3969	B
IM-11	2411	A	IM-47	4649	A	IM-83	132976	A	IM-129	3986	B	IM-130	3997	B
IM-12	2434	A	IM-48	4798	A	IM-84	149099	A	IM-131	4013	B	IM-132	4181	B
IM-13	2467	A	IM-49	5104	A	IM-85	2016	B	IM-133	4297	B	IM-134	4311	B
IM-14	2482	A	IM-50	5918	A	IM-86	2029	B	IM-135	4465	B	IM-136	4484	B
IM-15	2498	A	IM-51	6122	A	IM-87	2144	B	IM-137	4579	B	IM-138	4608	B
IM-16	2565	A	IM-52	6192	A	IM-88	2130	B	IM-139	4669	B	IM-140	4747	B
IM-17	2574	A	IM-53	6452	A	IM-89	2168	B	IM-141	4862	B	IM-142	4891	B
IM-18	2586	A	IM-54	6660	A	IM-90	2184	B	IM-143	5033	B	IM-144	5077	B
IM-19	2605	A	IM-55	7766	A	IM-91	2200	B						
IM-20	2722	A	IM-56	8145	A	IM-92	2284	B						
IM-21	2746	A	IM-57	8954	A	IM-93	2299	B						
IM-22	2788	A	IM-58	9312	A	IM-94	2314	B						
IM-23	2855	A	IM-59	9449	A	IM-95	2414	B						
IM-24	2871	A	IM-60	10272	A	IM-96	2428	B						
IM-25	2984	A	IM-61	11663	A	IM-97	2451	B						
IM-26	3030	A	IM-62	13376	A	IM-98	2466	B						
IM-27	3144	A	IM-63	14698	A	IM-99	2483	B						
IM-28	3243	A	IM-64	15190	A	IM-100	2565	B						
IM-29	3273	A	IM-64	66758	A	IM-101	2583	B						
IM-30	3290	A	IM-65	15951	A	IM-102	2597	B						
IM-31	3369	A	IM-66	15172	A	IM-103	2697	B						
IM-32	3445	A	IM-67	15925	A	IM-104	2715	B						
IM-33	3483	A	IM-68	23436	A	IM-105	2740	B						
IM-34	3676	A	IM-69	39794	A	IM-106	2752	B						
IM-35	3779	A	IM-70	44166	A	IM-107	2767	B						
IM-36	3793	A	IM-71	46890	A	IM-108	2865	B						

FIGURE 1B

MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION
IM-145	5099	B	IM-181	16018	B	IM-217	2130	C	IM-253	3733	C	IM-289	3733	C	IM-325	3733	C
IM-146	5143	B	IM-182	17346	B	IM-218	2145	C	IM-254	3833	C	IM-290	3833	C	IM-326	3833	C
IM-147	5158	B	IM-183	18311	B	IM-219	2167	C	IM-255	3900	C	IM-291	3900	C	IM-327	3900	C
IM-148	5272	B	IM-184	22586	B	IM-220	2182	C	IM-256	4010	C	IM-292	4010	C	IM-328	4010	C
IM-149	5306	B	IM-185	23422	B	IM-221	2199	C	IM-257	4145	C	IM-293	4145	C	IM-329	4145	C
IM-150	5349	B	IM-186	27969	B	IM-222	2211	C	IM-258	4187	C	IM-294	4187	C	IM-330	4187	C
IM-151	5364	B	IM-187	33283	B	IM-223	2230	C	IM-259	4299	C	IM-295	4299	C	IM-331	4299	C
IM-152	5421	B	IM-188	39808	B	IM-224	2250	C	IM-260	4466	C	IM-296	4466	C	IM-332	4466	C
IM-153	5554	B	IM-189	43110	B	IM-225	2280	C	IM-261	4582	C	IM-297	4582	C	IM-333	4582	C
IM-154	5711	B	IM-190	44454	B	IM-226	2297	C	IM-262	4813	C	IM-298	4813	C	IM-334	4813	C
IM-155	5876	B	IM-191	47215	B	IM-227	2317	C	IM-263	4876	C	IM-299	4876	C	IM-335	4876	C
IM-156	5916	B	IM-192	53784	B	IM-228	2412	C	IM-264	5032	C	IM-300	5032	C	IM-336	5032	C
IM-157	5931	B	IM-193	55952	B	IM-229	2428	C	IM-265	5347	C	IM-301	5347	C	IM-337	5347	C
IM-158	5988	B	IM-194	60573	B	IM-230	2468	C	IM-266	5365	C	IM-302	5365	C	IM-338	5365	C
IM-159	6137	B	IM-195	66346	B	IM-231	2481	C	IM-267	5932	C	IM-303	5932	C	IM-339	5932	C
IM-160	6200	B	IM-196	73387	B	IM-232	2498	C	IM-268	7767	C	IM-304	7767	C	IM-340	7767	C
IM-161	6443	B	IM-197	79203	B	IM-233	2567	C	IM-269	7973	C	IM-305	7973	C	IM-341	7973	C
IM-162	6644	B	IM-198	89302	B	IM-234	2585	C	IM-270	8143	C	IM-306	8143	C	IM-342	8143	C
IM-163	6958	B	IM-199	94226	B	IM-235	2599	C	IM-271	9187	C	IM-307	9187	C	IM-343	9187	C
IM-164	7481	B	IM-200	99358	B	IM-236	2698	C	IM-272	9293	C	IM-308	9293	C	IM-344	9293	C
IM-165	7568	B	IM-201	102096	B	IM-237	2715	C	IM-273	11705	C	IM-309	11705	C	IM-345	11705	C
IM-166	7765	B	IM-202	107199	B	IM-238	2745	C	IM-274	14041	C	IM-310	14041	C	IM-346	14041	C
IM-167	7955	B	IM-203	116936	B	IM-239	2766	C	IM-275	15114	C	IM-311	15114	C	IM-347	15114	C
IM-168	8144	B	IM-204	119487	B	IM-240	2867	C	IM-276	15939	C	IM-312	15939	C	IM-348	15939	C
IM-169	8698	B	IM-205	122103	B	IM-241	2885	C	IM-277	22321	C	IM-313	22321	C	IM-349	22321	C
IM-170	8821	B	IM-206	125431	B	IM-242	2998	C	IM-278	28001	C	IM-314	28001	C	IM-350	28001	C
IM-171	8944	B	IM-207	132052	B	IM-243	3052	C	IM-279	33296	C	IM-315	33296	C	IM-351	33296	C
IM-172	9138	B	IM-208	138518	B	IM-244	3096	C	IM-280	39770	C	IM-316	39770	C	IM-352	39770	C
IM-173	9298	B	IM-209	145147	B	IM-245	3151	C	IM-281	44460	C	IM-317	44460	C	IM-353	44460	C
IM-174	9390	B	IM-210	157502	B	IM-246	3167	C	IM-282	47307	C	IM-318	47307	C	IM-354	47307	C
IM-175	9516	B	IM-211	168579	B	IM-247	3286	C	IM-283	50625	C	IM-319	50625	C	IM-355	50625	C
IM-176	11711	B	IM-212	173391	B	IM-248	3303	C	IM-284	55898	C	IM-320	55898	C	IM-356	55898	C
IM-177	11914	B	IM-213	2011	C	IM-249	3335	C	IM-285	60882	C	IM-321	60882	C	IM-357	60882	C
IM-178	14033	B	IM-214	2030	C	IM-250	3448	C	IM-286	66294	C	IM-322	66294	C	IM-358	66294	C
IM-179	15110	B	IM-215	2050	C	IM-251	3619	C	IM-287	78892	C	IM-323	78892	C	IM-359	78892	C
IM-180	15838	B	IM-216	2096	C	IM-252	3709	C	IM-288	83848	C	IM-324	83848	C	IM-360	83848	C

MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION
IM-289	89081	C	IM-325	2565	D	IM-361	13857	D	IM-397	2082	E			
IM-290	94147	C	IM-326	2582	D	IM-362	14056	D	IM-398	2128	E			
IM-291	99324	C	IM-327	2597	D	IM-363	15108	D	IM-399	2148	E			
IM-292	107163	C	IM-328	2716	D	IM-364	15844	D	IM-400	2170	E			
IM-293	110350	C	IM-329	2747	D	IM-365	22243	D	IM-401	2187	E			
IM-294	113339	C	IM-330	2767	D	IM-366	22465	D	IM-402	2206	E			
IM-295	116291	C	IM-331	2866	D	IM-367	28022	D	IM-403	2232	E			
IM-296	122769	C	IM-332	2882	D	IM-368	33272	D	IM-404	2250	E			
IM-297	131908	C	IM-333	2994	D	IM-369	40149	D	IM-405	2279	E			
IM-298	145248	C	IM-334	3032	D	IM-370	43113	D	IM-406	2296	E			
IM-299	159252	C	IM-335	3050	D	IM-371	44219	D	IM-407	2314	E			
IM-300	165164	C	IM-336	3148	D	IM-372	47196	D	IM-408	2354	E			
IM-301	174928	C	IM-337	3164	D	IM-373	51062	D	IM-409	2394	E			
IM-302	196003	C	IM-338	3278	D	IM-374	56082	D	IM-410	2413	E			
IM-303	2007	D	IM-339	3334	D	IM-375	58239	D	IM-411	2436	E			
IM-304	2016	D	IM-340	3385	D	IM-376	60285	D	IM-412	2457	E			
IM-305	2030	D	IM-341	3432	D	IM-377	66148	D	IM-413	2466	E			
IM-306	2052	D	IM-342	3451	D	IM-378	73668	D	IM-414	2499	E			
IM-307	2099	D	IM-343	3617	D	IM-379	77433	D	IM-415	2566	E			
IM-308	2130	D	IM-344	3701	D	IM-380	79986	D	IM-416	2583	E			
IM-309	2144	D	IM-345	3725	D	IM-381	80844	D	IM-417	2612	E			
IM-310	2154	D	IM-346	3833	D	IM-382	88962	D	IM-418	2662	E			
IM-311	2166	D	IM-347	3899	D	IM-383	94399	D	IM-419	2723	E			
IM-312	2184	D	IM-348	4008	D	IM-384	99419	D	IM-420	2738	E			
IM-313	2204	D	IM-349	4157	D	IM-385	108395	D	IM-421	2750	E			
IM-314	2231	D	IM-350	4297	D	IM-386	116433	D	IM-422	2849	E			
IM-315	2252	D	IM-351	4580	D	IM-387	123337	D	IM-423	2867	E			
IM-316	2275	D	IM-352	4805	D	IM-388	131977	D	IM-424	3036	E			
IM-317	2299	D	IM-353	6946	D	IM-389	145658	D	IM-425	3147	E			
IM-318	2316	D	IM-354	7053	D	IM-390	152603	D	IM-426	3281	E			
IM-319	2412	D	IM-355	7767	D	IM-391	159524	D	IM-427	3319	E			
IM-320	2435	D	IM-356	7954	D	IM-392	196072	D	IM-428	3445	E			
IM-321	2466	D	IM-357	8139	D	IM-393	2010	E	IM-429	3693	E			
IM-322	2480	D	IM-358	9292	D	IM-394	2029	E	IM-430	3731	E			
IM-323	2499	D	IM-359	11671	D	IM-395	2050	E	IM-431	3818	E			
IM-324	2518	D	IM-360	13727	D	IM-396	2068	E	IM-432	3885	E			

FIGURE 1D

MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION	MARKER ID	MW	FRACTION
IM-433	4136	E	IM-469	86211	E	IM-505	4174	F	IM-541	95033	F	IM-541	95033	F
IM-434	4169	E	IM-470	89407	E	IM-506	4362	F	IM-542	100310	F	IM-542	100310	F
IM-435	4257	E	IM-471	100270	E	IM-507	4473	F	IM-543	116889	F	IM-543	116889	F
IM-436	4277	E	IM-472	109638	E	IM-508	4631	F	IM-544	132711	F	IM-544	132711	F
IM-437	4355	E	IM-473	117132	E	IM-509	4822	F	IM-545	147276	F	IM-545	147276	F
IM-438	4369	E	IM-474	132843	E	IM-510	5862	F	IM-546	160768	F	IM-546	160768	F
IM-439	4470	E	IM-475	147160	E	IM-511	6192	F						
IM-440	4486	E	IM-476	152199	E	IM-512	6941	F						
IM-441	4541	E	IM-477	166461	E	IM-513	7626	F						
IM-442	4634	E	IM-478	176635	E	IM-514	7772	F						
IM-443	4841	E	IM-479	2011	F	IM-515	7957	F						
IM-444	5862	E	IM-480	2030	F	IM-516	8150	F						
IM-445	5911	E	IM-481	2128	F	IM-517	8954	F						
IM-446	6649	E	IM-482	2149	F	IM-518	9300	F						
IM-447	6952	E	IM-483	2186	F	IM-519	11545	F						
IM-448	7769	E	IM-484	2207	F	IM-520	11717	F						
IM-449	8148	E	IM-485	2279	F	IM-521	13887	F						
IM-450	8260	E	IM-486	2299	F	IM-522	14073	F						
IM-451	8785	E	IM-487	2319	F	IM-523	15196	F						
IM-452	9301	E	IM-488	2412	F	IM-524	15903	F						
IM-453	10071	E	IM-489	2434	F	IM-525	22460	F						
IM-454	11721	E	IM-490	2467	F	IM-526	23135	F						
IM-455	13910	E	IM-491	2485	F	IM-527	28135	F						
IM-456	15919	E	IM-492	2582	F	IM-528	33577	F						
IM-457	22422	E	IM-493	2605	F	IM-529	39813	F						
IM-458	28233	E	IM-494	2697	F	IM-530	42344	F						
IM-459	33490	E	IM-495	2751	F	IM-531	43274	F						
IM-460	43121	E	IM-496	2865	F	IM-532	44345	F						
IM-461	44558	E	IM-497	3036	F	IM-533	51007	F						
IM-462	46694	E	IM-498	3151	F	IM-534	56318	F						
IM-463	50954	E	IM-499	3372	F	IM-535	60079	F						
IM-464	54478	E	IM-500	3440	F	IM-536	66690	F						
IM-465	60041	E	IM-501	3488	F	IM-537	75122	F						
IM-466	66652	E	IM-502	3717	F	IM-538	78429	F						
IM-467	75580	E	IM-503	3890	F	IM-539	81249	F						
IM-468	79463	E	IM-504	4155	F	IM-540	89384	F						

FIGURE 2

RANK	MW	MARKER ID	RANK	MW	MARKER ID
1	14073	IM-522	39	117132	IM-473
2	11705	IM-273	40	13727	IM-360
3	11717	IM-520	41	4257	IM-435
4	11545	IM-519	42	5349	IM-150
5	11721	IM-454	43	5364	IM-151
6	4473	IM-507	44	2967	IM-110
7	13887	IM-521	45	6122	IM-51
8	5272	IM-148	46	6958	IM-163
9	5365	IM-266	47	4355	IM-437
10	75122	IM-537	48	160768	IM-546
11	100270	IM-471	49	5554	IM-153
12	5862	IM-510	50	7767	IM-268
13	132711	IM-544			
14	132843	IM-474			
15	5876	IM-155			
16	5932	IM-157			
17	11711	IM-176			
18	5911	IM-445			
19	11914	IM-177			
20	4486	IM-440			
21	79463	IM-468			
22	4369	IM-438			
23	100310	IM-542			
24	11671	IM-359			
25	4277	IM-436			
26	2752	IM-106			
27	13910	IM-455			
28	5862	IM-444			
29	5988	IM-158			
30	5347	IM-265			
31	5918	IM-50			
32	6137	IM-159			
33	5916	IM-156			
34	4470	IM-439			
35	5931	IM-157			
36	4631	IM-508			
37	7772	IM-514			
38	176635	IM-478			

Figure 3A

[illegible]

Figure 3c

[illegible]

78755	78756	78757	78758	78759	78760	78804	78805	78916	78917	78918	78919	78920	78921	78922	78923	78924	78925	78926	78927	78928	78929	78930	78931	78932	78933	78934	78935	78936	78937	78938	78939	78940	78941	78942	78943	78944	78945	78946	78947	78948	78949	78950	78951	78952	78953	78954	78955	78956	78957	78958	78959	78960	78961	78962	78963	78964	78965	78966	78967	78968	78969	78970	78971	78972	78973	78974	78975	78976	78977	78978	78979	78980	78981	78982	78983	78984	78985	78986	78987	78988	78989	78990	78991	78992	78993	78994	78995	78996	78997	78998	78999	79000	79001	79002	79003	79004	79005	79006	79007	79008	79009	79010	79011	79012	79013	79014	79015	79016	79017	79018	79019	79020	79021	79022	79023	79024	79025	79026	79027	79028	79029	79030	79031	79032	79033	79034	79035	79036	79037	79038	79039	79040	79041	79042	79043	79044	79045	79046	79047	79048	79049	79050	79051	79052	79053	79054	79055	79056	79057	79058	79059	79060	79061	79062	79063	79064	79065	79066	79067	79068	79069	79070	79071	79072	79073	79074	79075	79076	79077	79078	79079	79080	79081	79082	79083	79084	79085	79086	79087	79088	79089	79090	79091	79092	79093	79094	79095	79096	79097	79098	79099	79100	79101	79102	79103	79104	79105	79106	79107	79108	79109	79110	79111	79112	79113	79114	79115	79116	79117	79118	79119	79120	79121	79122	79123	79124	79125	79126	79127	79128	79129	79130	79131	79132	79133	79134	79135	79136	79137	79138	79139	79140	79141	79142	79143	79144	79145	79146	79147	79148	79149	79150	79151	79152	79153	79154	79155	79156	79157	79158	79159	79160	79161	79162	79163	79164	79165	79166	79167	79168	79169	79170	79171	79172	79173	79174	79175	79176	79177	79178	79179	79180	79181	79182	79183	79184	79185	79186	79187	79188	79189	79190	79191	79192	79193	79194	79195	79196	79197	79198	79199	79200	79201	79202	79203	79204	79205	79206	79207	79208	79209	79210	79211	79212	79213	79214	79215	79216	79217	79218	79219	79220	79221	79222	79223	79224	79225	79226	79227	79228	79229	79230	79231	79232	79233	79234	79235	79236	79237	79238	79239	79240	79241	79242	79243	79244	79245	79246	79247	79248	79249	79250	79251	79252	79253	79254	79255	79256	79257	79258	79259	79260	79261	79262	79263	79264	79265	79266	79267	79268	79269	79270	79271	79272	79273	79274	79275	79276	79277	79278	79279	79280	79281	79282	79283	79284	79285	79286	79287	79288	79289	79290	79291	79292	79293	79294	79295	79296	79297	79298	79299	79300	79301	79302	79303	79304	79305	79306	79307	79308	79309	79310	79311	79312	79313	79314	79315	79316</
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WM-204	C	14093	14098	14078	14086	14088	14086	14088	14096
WM-205	C	15148	15141	15138	15130	15139	15139	15139	15139
WM-206	C	15882	15870	15882	15854	15887	15887	15885	15885
WM-207	C	17365	17367	17367	17359	17369	17367	17367	17367
WM-208	C	22258	22250	22258	22252	22260	22248	22261	22268
WM-209	C	28110	28117	28123	28120	28126	28110	28114	28129
WM-210	C	33310	33316	33323	33316	33320	33314	33312	33339
WM-211	C	37250	37214	37278	37238	37236	37236	37236	37270
WM-212	C	44528	44528	44513	44444	44444	44542	44518	44518
WM-213	C	47378	47378	47380	47391	47384	47377	47377	47423
WM-214	C	51210	51280	51284	51297	51297	51237	51369	51268
WM-215	C	56204	56230	56240	56176	56212	56296	56536	56536
WM-216	C	60781	59763	59764	59695	59710	59758	59752	60781
WM-217	C	65443	65443	65443	65391	65391	65443	65447	65447
WM-218	C	74748	74514	74695	74701	74780	74829	74829	74587
WM-219	C	78519	78519	78738	78655	78655	78608	78651	78948
WM-220	C	80465	80465	80375	80360	80360	80557	80772	80468
WM-221	C	83659	83659	83617	83674	83674	83612	83612	83436
WM-222	C	88528	88528	88581	88588	88588	88514	88590	89950
WM-223	C	94716	94716	94672	94624	94624	94725	94711	94738
WM-224	C	99819	99819	99739	99895	99895	99738	99763	99798
WM-225	C	110775	110758	110804	110825	110825	110629	110738	110815
WM-226	C	116193	116193	116238	116238	116238	116237	116098	116344
WM-227	C	132423	132397	132417	132434	132426	132417	132388	132435
WM-228	C	145302	145302	145361	145361	145361	145477	145617	145617
WM-229	C	154931	154931	154931	154931	154930	155075	155082	155075
WM-230	C	165378	165378	165369	165402	165402	165431	165374	165374
WM-231	C	177360	177164	177208	177355	177355	177058	177024	176877
WM-232	C	181213	181213	181213	181213	181213	181213	181213	181213
WM-233	C	198220	198220	198220	198220	198220	198220	198220	198220
WM-234	D	2011	2010	2010	2009	2009	2010	2010	2010
WM-235	D	2032	2032	2032	2032	2032	2032	2031	2033
WM-236	D	2054	2054	2054	2054	2054	2053	2053	2054
WM-237	D	2078	2078	2078	2078	2078	2078	2078	2078
WM-238	D	2167	2167	2167	2167	2167	2168	2168	2167
WM-239	D	2185	2185	2185	2185	2185	2185	2185	2185
WM-240	D	2211	2212	2211	2210	2210	2211	2211	2212
WM-241	D	2230	2230	2231	2228	2228	2229	2229	2229
WM-242	D	2258	2258	2258	2254	2254	2255	2255	2256
WM-243	D	2278	2277	2277	2278	2278	2278	2278	2278
WM-244	D	2299	2299	2300	2298	2298	2299	2299	2299
WM-245	D	2354	2354	2354	2352	2352	2352	2353	2354
WM-246	D	2427	2428	2428	2427	2427	2427	2427	2428
WM-247	D	2481	2481	2483	2478	2478	2480	2481	2481
WM-248	D	2500	2500	2500	2500	2500	2500	2500	2500
WM-249	D	2538	2538	2531	2543	2543	2542	2542	2499
WM-250	D	2578	2578	2578	2578	2578	2542	2542	2578
WM-251	D	2637	2637	2637	2637	2637	2637	2637	2637
WM-252	D	2751	2751	2754	2748	2748	2748	2750	2750
WM-253	D	2865	2865	2865	2867	2867	2864	2864	2864
WM-254	D	2937	2937	2938	2938	2938	2936	2936	2936

Figure 3F

WM-255	D	3081	3082	3085	3086	3087	3088	3089	3093
WM-256	D	3153	3158	3159	3159	3157	3157	3157	3154
WM-257	D	3281	3282	3282	3283	3283	3285	3283	3282
WM-258	D	3376	3376	3366	3366	3370	3370	3370	3365
WM-259	D	3445	3448	3445	3445	3443	3442	3446	3444
WM-260	D	3559	3556	3558	3556	3557	3557	3557	3557
WM-261	D	3685	3688	3688	3688	3684	3683	3688	3686
WM-262	D	3815	3815	3818	3813	3814	3813	3816	3814
WM-263	D	3943	3942	3943	3939	3941	3941	3942	3942
WM-264	D	3957	3953	3953	3949	3951	3950	3951	3950
WM-265	D	4052	4052	4052	4052	4052	4052	4051	4150
WM-266	D	4130	4132	4131	4128	4129	4129	4129	4208
WM-267	D	4210	4207	4202	4207	4206	4208	4208	4410
WM-268	D	4800	4795	4801	4768	4801	4767	4768	4765
WM-269	D	6438	6437	6435	6432	6435	6437	6437	6437
WM-270	D	6648	6629	6635	6632	6637	6637	6638	6680
WM-271	D	6859	6848	6847	6841	6844	6861	6851	6948
WM-272	D	7573	7574	7571	7571	7574	7574	7572	7573
WM-273	D	7776	7760	7762	7762	7771	7767	7768	7764
WM-274	D	7942	7943	7943	7942	7944	7944	7948	7947
WM-275	D	11505	11498	11497	11498	11498	11498	11500	11497
WM-276	D	11678	11678	11673	11682	11678	11683	11683	11678
WM-277	D	13777	13778	13777	13771	13776	13776	13774	13773
WM-278	D	13683	13686	13685	13671	13675	13689	13689	13692
WM-279	D	15142	15136	15144	15128	15134	15138	15137	15140
WM-280	D	15876	15874	15874	15863	15871	15881	15885	15888
WM-281	D	17347	17322	17350	17341	17346	17341	17346	17346
WM-282	D	22275	22277	22275	22274	22277	22288	22287	22287
WM-283	D	28121	28135	28129	28130	28119	28139	28135	28135
WM-284	D	33333	33364	33320	33347	33336	33365	33367	33367
WM-285	D	37251	37381	37271	37306	37306	37322	37276	37232
WM-286	D	40281	40259	40304	40255	40261	40266	40266	40266
WM-287	D	44131	44135	44133	44502	44518	44558	44558	44558
WM-288	D	44513	44493	44549	44502	44518	44558	44558	44558
WM-289	D	51234	51228	51234	51224	51209	51235	51238	51262
WM-290	D	55648	55659	55645	55654	55614	55658	55655	55667
WM-291	D	66435	66437	66404	66470	66404	66469	66468	66469
WM-292	D	73586	73753	73767	73587	73587	73654	73775	73686
WM-293	D	75088	78088	75064	75060	75133	78098	78098	78103
WM-294	D	80410	80224	80224	80234	80226	80448	80448	80448
WM-295	D	80305	81643	81600	81659	81659	81659	81659	81659
WM-296	D	88834	88990	88849	88857	88875	88967	88940	88940
WM-297	D	94507	94554	94451	94420	94520	94540	94514	94543

Figure 3G

154336	160431	160716	160873	153878	160722	160347	160311
154337	160432	160717	160874	153879	160723	160348	160312
154338	160433	160718	160875	153880	160724	160349	160313
154339	160434	160719	160876	153881	160725	160350	160314
154340	160435	160720	160877	153882	160726	160351	160315
154341	160436	160721	160878	153883	160727	160352	160316
154342	160437	160722	160879	153884	160728	160353	160317
154343	160438	160723	160880	153885	160729	160354	160318
154344	160439	160724	160881	153886	160730	160355	160319
154345	160440	160725	160882	153887	160731	160356	160320
154346	160441	160726	160883	153888	160732	160357	160321
154347	160442	160727	160884	153889	160733	160358	160322
154348	160443	160728	160885	153890	160734	160359	160323
154349	160444	160729	160886	153891	160735	160360	160324
154350	160445	160730	160887	153892	160736	160361	160325
154351	160446	160731	160888	153893	160737	160362	160326
154352	160447	160732	160889	153894	160738	160363	160327
154353	160448	160733	160890	153895	160739	160364	160328
154354	160449	160734	160891	153896	160740	160365	160329
154355	160450	160735	160892	153897	160741	160366	160330
154356	160451	160736	160893	153898	160742	160367	160331
154357	160452	160737	160894	153899	160743	160368	160332
154358	160453	160738	160895	153900	160744	160369	160333
154359	160454	160739	160896	153901	160745	160370	160334
154360	160455	160740	160897	153902	160746	160371	160335
154361	160456	160741	160898	153903	160747	160372	160336
154362	160457	160742	160899	153904	160748	160373	160337
154363	160458	160743	160900	153905	160749	160374	160338
154364	160459	160744	160901	153906	160750	160375	160339
154365	160460	160745	160902	153907	160751	160376	160340
154366	160461	160746	160903	153908	160752	160377	160341
154367	160462	160747	160904	153909	160753	160378	160342
154368	160463	160748	160905	153910	160754	160379	160343
154369	160464	160749	160906	153911	160755	160380	160344
154370	160465	160750	160907	153912	160756	160381	160345
154371	160466	160751	160908	153913	160757	160382	160346
154372	160467	160752	160909	153914	160758	160383	160347
154373	160468	160753	160910	153915	160759	160384	160348
154374	160469	160754	160911	153916	160760	160385	160349
154375	160470	160755	160912	153917	160761	160386	160350
154376	160471	160756	160913	153918	160762	160387	160351
154377	160472	160757	160914	153919	160763	160388	160352
154378	160473	160758	160915	153920	160764	160389	160353
154379	160474	160759	160916	153921	160765	160390	160354
154380	160475	160760	160917	153922	160766	160391	160355
154381	160476	160761	160918	153923	160767	160392	160356
154382	160477	160762	160919	153924	160768	160393	160357
154383	160478	160763	160920	153925	160769	160394	160358
154384	160479	160764	160921	153926	160770	160395	160359
154385	160480	160765	160922	153927	160771	160396	160360
154386	160481	160766	160923	153928	160772	160397	160361
154387	160482	160767	160924	153929	160773	160398	160362
154388	160483	160768	160925	153930	160774	160399	160363
154389	160484	160769	160926	153931	160775	160400	160364
154390	160485	160770	160927	153932	160776	160401	160365
154391	160486	160771	160928	153933	160777	160402	160366
154392	160487	160772	160929	153934	160778	160403	160367
154393	160488	160773	160930	153935	160779	160404	160368
154394	160489	160774	160931	153936	160780	160405	160369
154395	160490	160775	160932	153937	160781	160406	160370
154396	160491	160776	160933	153938	160782	160407	160371
154397	160492	160777	160934	153939	160783	160408	160372
154398	160493	160778	160935	153940	160784	160409	160373
154399	160494	160779	160936	153941	160785	160410	160374
154400	160495	160780	160937	153942	160786	160411	160375
154401	160496	160781	160938	153943	160787	160412	160376
154402	160497	160782	160939	153944	160788	160413	160377
154403	160498	160783	160940	153945	160789	160414	160378
154404	160499	160784	160941	153946	160790	160415	160379
154405	160500	160785	160942	153947	160791	160416	160380
154406	160501	160786	160943	153948	160792	160417	160381
154407	160502	160787	160944	153949	160793	160418	160382
154408	160503	160788	160945	153950	160794	160419	160383
154409	160504	160789	160946	153951	160795	160420	160384
154410	160505	160790	160947	153952	160796	160421	160385
154411	160506	160791	160948	153953	160797	160422	160386
154412	160507	160792	160949	153954	160798	160423	160387
154413	160508	160793	160950	153955	160799	160424	160388
154414	160509	160794	160951	153956	160800	160425	160389
154415	160510	160795	160952	153957	160801	160426	160390
154416	160511	160796	160953	153958	160802	160427	160391
154417	160512	160797	160954	153959	160803	160428	160392
154418	160513	160798	160955	153960	160804	160429	160393
154419	160514	160799	160956	153961	160805	160430	160394
154420	160515	160800	160957	153962	160806	160431	160395
154421	160516	160801	160958	153963	160807	160432	160396
154422	160517	160802	160959	153964	160808	160433	160397
154423	160518	160803	160960	153965	160809	160434	160398
154424	160519	160804	160961	153966	160810	160435	160399
154425	160520	160805	160962	153967	160811	160436	160400
154426	160521	160806	160963	153968	160812	160437	160401
154427	160522	160807	160964	153969	160813	160438	160402
154428	160523	160808	160965	153970	160814	160439	160403
154429	160524	160809	160966	153971	160815	160440	160404
154430	160525	160810	160967	153972	160816	160441	160405
154431	160526	160811	160968	153973	160817	160442	160406
154432	160527	160812	160969	153974	160818	160443	160407
154433	160528	160813	160970	153975	160819	160444	160408
154434	160529	160814	160971	153976	160820	160445	160409
154435	160530	160815	160972	153977	160821	160446	160410
154436	160531	160816	160973	153978	160822	160447	160411
154437	160532	160817	160974	153979	160823	160448	160412
154438	160533	160818	160975	153980	160824	160449	160413
154439	160534	160819	160976	153981	160825	160450	160414
154440	160535	160820	160977	153982	160826	160451	160415
154441	160536	160821	160978	153983	160827	160452	160416
154442	160537	160822	160979	153984	160828	160453	160417
154443	160538	160823	160980	153985	160829	160454	160418
154444	160539	160824	160981	153986	160830	160455	160419
154445	160540	160825	160982	153987	160831	160456	160420
154446	160541	160826	160983	153988	160832	160457	160421
154447	160542	160827	160984	153989	160833	160458	160422
154448	160543	160828	160985	153990	160834	160459	160423
154449	160544	160829	160986	153991	160835	160460	160424
154450	160545	160830	160987	153992	160836	160461	160425
154451	160546	160831	160988	153993	160837	160462	160426
154452	160547	160832	160989	153994	160838	160463	160427
154453	160548	160833	160990	153995	160839	160464	160428
154454	160549	160834	160991	153996	160840	160465	160429
154455	160550	160835	160992	153997	160841	160466	160430
154456	160551	160836	160993	153998	160842	160467	160431
154457	160552	160837	160994	153999	160843	160468	160432
154458	160553	160838	160995	154000	160844	160469	160433
154459	160554	160839	160996	154001	160845	160470	160434
154460	160555	160840	160997	154002	160846	160471	160435
154461	160556	160841	160998	154003	160847	160472	160436
154462	160557	160842	160999	154004	160848	160473	160437
154463	160558	160843	161000	154005	160849	160474	160438
154464	160559	160844	161001	154006	160850	160475	160439
154465	160560	160845	161002	154007	160851	160476	160440
154466	160561	160846	161003	154008	160852	160477	160441
154467	160562	160847	161004	154009	160853	160478	160442
154468	160563	160848	161005	154010	160854	160479	160443
154469	160564	160849	161006	154011	160855	160480	160444
154470	160565	160850	161007	154012			

WM-357	E	8628	8854	8628	8827	8631	8630
WM-358	E	8849	8955	8955	8952	8945	8946
WM-359	E	8176	9181	9176	9177	9177	9194
WM-360	E	9302	9287	9289	9301	9301	9286
WM-361	E	9470	9478	9454	9452	9470	9477
WM-362	E	11530	11532	11515	11520	11533	11527
WM-363	E	11714	11699	11687	11687	11716	11716
WM-364	E	12482	12450	12464	12452	12461	12460
WM-365	E	12814	12804	12813	12822	12817	12818
WM-366	E	13897	13876	13888	13871	13901	13907
WM-367	E	15148	15150	15154	15154	15165	15160
WM-368	E	15891	15891	15901	15899	15899	15896
WM-369	E	17354	17370	17355	17362	17363	17365
WM-370	E	22368	22345	22348	22322	22384	22332
WM-371	E	28169	28163	28162	28165	28163	28172
WM-372	E	33429	33451	33431	33428	33425	33413
WM-373	E	39455	39235				39238
WM-374	E	44803	44596	44627	44591	44608	44651
WM-375	E	51376	51359	51378	51370	51400	51409
WM-376	E	59798	59842	59828	59847	59799	59799
WM-377	E	66557	66553	66558	66527	66581	66504
WM-378	E	75345	75449	75345	75201	75488	75419
WM-379	E	78354	78334	78388	78858	78390	78327
WM-380	E	88411	88288	88400	88370	88421	88542
WM-381	E	99991	99991	98979	98992	100148	100028
WM-382	E	110157	109554	109594	109916	110474	110414
WM-383	E	117763	117603	117760	117739	117699	117802
WM-384	E	132840	132828	132845	132827	132853	132842
WM-385	E	146531	146537	146553	146208	145912	146953
WM-386	E	156171	148507			152548	152534
WM-387	E	165715	165723	165192	165727	165485	165575
WM-388	E	182879	182854	182289	182217	183093	183596
WM-389	E	197328	197348	197208	197264	197248	197301
WM-390	F	2011	2010	2010	2012	2011	2011
WM-391	F	2031	2029	2029	2030	2029	2029
WM-392	F	2055	2049	2049	2053	2052	2052
WM-393	F	2085	2088	2087	2071	2070	2067
WM-394	F	2085	2083	2083	2084	2084	2080
WM-395	F	2128	2128	2126	2126	2126	2126
WM-396	F	2186	2167	2167	2167	2167	2166
WM-397	F	2214	2213	2214	2212	2213	2213
WM-398	F	2234	2234	2234	2234	2234	2234
WM-399	F	2280	2278	2277	2280	2278	2278
WM-400	F	2298	2298	2297	2297	2297	2296
WM-401	F	2388	2390	2389	2389	2389	2389
WM-402	F	2413	2413	2412	2413	2413	2412
WM-403	F	2482	2482	2479	2474	2482	2481
WM-404	F	2500	2500	2500	2501	2500	2500
WM-405	F	2569	2569	2570	2569	2568	2568
WM-406	F	2581	2581	2582	2581	2581	2582
WM-407	F	2697	2697	2700	2695	2696	2696

Figure 3I

WN-408	F	2722	2723	2724	2725	2726	2727	2728	2729	2730	2731	2732	2733	2734	2735	2736	2737	2738	2739	2740	2741	2742	2743	2744	2745	2746	2747	2748	2749	2750	2751	2752	2753	2754	2755	2756	2757	2758	2759	2760	2761	2762	2763	2764	2765	2766	2767	2768	2769	2770	2771	2772	2773	2774	2775	2776	2777	2778	2779	2780	2781	2782	2783	2784	2785	2786	2787	2788	2789	2790	2791	2792	2793	2794	2795	2796	2797	2798	2799	2800	2801	2802	2803	2804	2805	2806	2807	2808	2809	2810	2811	2812	2813	2814	2815	2816	2817	2818	2819	2820	2821	2822	2823	2824	2825	2826	2827	2828	2829	2830	2831	2832	2833	2834	2835	2836	2837	2838	2839	2840	2841	2842	2843	2844	2845	2846	2847	2848	2849	2850	2851	2852	2853	2854	2855	2856	2857	2858	2859	2860	2861	2862	2863	2864	2865	2866	2867	2868	2869	2870	2871	2872	2873	2874	2875	2876	2877	2878	2879	2880	2881	2882	2883	2884	2885	2886	2887	2888	2889	2890	2891	2892	2893	2894	2895	2896	2897	2898	2899	2900	2901	2902	2903	2904	2905	2906	2907	2908	2909	2910	2911	2912	2913	2914	2915	2916	2917	2918	2919	2920	2921	2922	2923	2924	2925	2926	2927	2928	2929	2930	2931	2932	2933	2934	2935	2936	2937	2938	2939	2940	2941	2942	2943	2944	2945	2946	2947	2948	2949	2950	2951	2952	2953	2954	2955	2956	2957	2958	2959	2960	2961	2962	2963	2964	2965	2966	2967	2968	2969	2970	2971	2972	2973	2974	2975	2976	2977	2978	2979	2980	2981	2982	2983	2984	2985	2986	2987	2988	2989	2990	2991	2992	2993	2994	2995	2996	2997	2998	2999	3000
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Figure 3J

WN-459	F	33478	33451	33504	33488	33497	33512	33487	33520
WN-460	F	43423	43430	43434	43393	43472	43898	43547	43416
WN-461	F	44565	44523	44758	44688	44678	44694	44678	44692
WN-462	F	51328	51176	52410	51183	51283	51228	51277	51052
WN-463	F	59467	59461	59444	59517	59508	59610	59468	59617
WN-464	F	59676	59805	59844	59495	59728	59656	59858	59470
WN-465	F	68615	68828	68808	68570	68832	68597	68674	68548
WN-466	F	75421	75416	75551	75313	75236	75464	75466	75438
WN-467	F		78308	78348	78184		78346	78278	78684
WN-468	F			83331			83307	83004	83132
WN-469	F	89778	88516	88751	89588	88549	89776	89740	89988
WN-470	F	94791	94707	94828	94765	94742	94628	94749	94653
WN-471	F	100416	100480	100814	100355	100386	100594	100524	100828
WN-472	F	117035	118878	117130	116883	117127	116803	116782	116883
WN-473	F	132765	133688	132940	132756	132728	132707	132913	132834
WN-474	F	148470	148725	148890	148682	148662	148500	148191	148210
WN-475	F	160334	161462		160783	160845	160342	160004	160570
WN-476	A			2184	2131	2128			
WN-477	A			2414		2139			
WN-478	A					2184			
WN-479	A					2415			
WN-480	A					2434			
WN-481	A					2451			
WN-482	A					2468			
WN-483	A	2587	2585	2568	2587	2568			
WN-484	A					2569			
WN-485	A					2568			
WN-486	A					2567			
WN-487	A					2567			
WN-488	A					2567			
WN-489	A					2567			
WN-490	A					2567			
WN-491	A					2567			
WN-492	A					2567			
WN-493	A					2567			
WN-494	A					2567			
WN-495	A					2567			
WN-496	A					2567			
WN-497	A					2567			
WN-498	A					2567			
WN-499	A					2567			
WN-500	A					2567			
WN-501	A					2567			
WN-502	A					2567			
WN-503	A					2567			
WN-504	A					2567			
WN-505	A					2567			
WN-506	A					2567			
WN-507	A					2567			
WN-508	A					2567			
WN-509	A					2567			

Figure 3k

WM-510 A				4753	
WM-511 A				4800	
WM-512 A				4878	
WM-513 A		5093		5097	5093
WM-514 A		5175		5175	
WM-515 A		5214		5211	
WM-516 A				5371	
WM-517 A				5572	
WM-518 A				5678	
WM-519 A		5997		6998	
WM-520 A	5999	6176		6187	
WM-521 A	6191			6480	
WM-522 A				6506	
WM-523 A				6841	
WM-524 A				7152	
WM-525 A				7192	
WM-526 A				7478	
WM-527 A		7481		7650	
WM-528 A	7487	7477		7829	
WM-529 A	7855	7929		8329	
WM-530 A		8331		8453	
WM-531 A	8333			9100	
WM-532 A				9183	
WM-533 A		9501		9811	
WM-534 A				11389	
WM-535 A	9518	9510		11640	
WM-536 A				11734	
WM-537 A				12718	
WM-538 A		13589		13592	
WM-539 A				14079	
WM-540 A	13593			16268	
WM-541 A	16280	16268		17079	
WM-542 A				17432	
WM-543 A				18391	
WM-544 A				32667	
WM-545 A				46710	
WM-546 A				111238	
WM-547 A				117809	
WM-548 A				133294	
WM-549 A				135741	
WM-550 A				176527	
WM-551 A				2038	
WM-552 A				2038	
WM-553 A				2182	
WM-554 A				2182	
WM-555 A				2212	
WM-556 B	2031	2031		2411	
WM-557 B	2183	2183		2449	
WM-558 B					
WM-559 B					
WM-560 B					

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Figure 3L

WM-581 B	2540	2539	2467	2467	2467
WM-582 B		2539	2539	2539	2539
WM-583 B	2978	2978	2978	2978	2978
WM-584 B	3181	3181	3182	3182	3182
WM-585 B		3274	3274	3274	3274
WM-586 B	3360	3360	3360	3360	3360
WM-587 B	3376	3376	3376	3376	3376
WM-588 B		3464	3464	3464	3464
WM-589 B					
WM-590 B	3768	3771	3855	3855	3855
WM-591 B			3770	3770	3770
WM-592 B			3842	3842	3842
WM-593 B					
WM-594 B			4080	4080	4080
WM-595 B	4092	4091	4092	4092	4092
WM-596 B	4148	4148	4147	4147	4147
WM-597 B	4647	4647	4647	4647	4647
WM-598 B	4790	4784	4793	4793	4792
WM-599 B		4855			
WM-600 B	5442	5443	5442	5442	5442
WM-601 B	5987	5988	5987	5987	5987
WM-602 B	6187	6184	6184	6184	6184
WM-603 B		6805			
WM-604 B			10558	10558	10558
WM-605 B		16057	17902	17902	17902
WM-606 B					
WM-607 B	83912	83843	83827	83827	83827
WM-608 B			53722	53722	53722
WM-609 B					
WM-610 B			159465	159465	159465
WM-611 B			171916	171916	171916
WM-612 B			2053	2053	2053
WM-613 B	2126	2126	2126	2126	2126
WM-614 B			2255	2255	2255
WM-615 B					
WM-616 B	2458	2457	2410	2410	2410
WM-617 B			2462	2462	2462
WM-618 B			2495	2495	2495
WM-619 B			2566	2566	2566
WM-620 B	2685	2689	2691	2691	2691
WM-621 B	2724	2889	2725	2725	2725
WM-622 B					
WM-623 B					
WM-624 B	2884	2982			
WM-625 B		3122			
WM-626 B			2679	2679	2679
WM-627 B	3163	3174	3162	3162	3162
WM-628 B	3178	3190	3177	3177	3177
WM-629 B		3252			
WM-630 B		3308			
WM-631 B					
WM-632 B					
WM-633 B					
WM-634 B					
WM-635 B					
WM-636 B					
WM-637 B					
WM-638 B					
WM-639 B					
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WM-664 B					
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WM-700 B					
WM-701 B					
WM-702 B					
WM-703 B					
WM-704 B					
WM-705 B					
WM-706 B					
WM-707 B					
WM-708 B					
WM-709 B					
WM-710 B					
WM-711 B					
WM-712 B					
WM-713 B					
WM-714 B					
WM-715 B					
WM-716 B					
WM-717 B					
WM-718 B					
WM-719 B					
WM-720 B					
WM-721 B					
WM-722 B					
WM-723 B					
WM-724 B					
WM-725 B					
WM-726 B					
WM-727 B					
WM-728 B					
WM-729 B					
WM-730 B					
WM-731 B					
WM-732 B					
WM-733 B					
WM-734 B					
WM-735 B					
WM-736 B					
WM-737 B					
WM-738 B					
WM-739 B					
WM-740 B					
WM-741 B					
WM-742 B					
WM-743 B					
WM-744 B					
WM-745 B					
WM-746 B					
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WM-748 B					
WM-749 B					
WM-750 B					
WM-751 B					
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WM-763 B					
WM-764 B					
WM-765 B					
WM-766 B					
WM-767 B					
WM-768 B					
WM-769 B					
WM-770 B					
WM-771 B					
WM-772 B					
WM-773 B					
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WM-775 B					
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WM-784 B					
WM-785 B					
WM-786 B					
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WM-790 B					
WM-791 B					
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WM-798 B					
WM-799 B					
WM-800 B					
WM-801 B					
WM-802 B					
WM-803 B					
WM-804 B					
WM-805 B					
WM-806 B					
WM-807 B					
WM-808 B					
WM-809 B					
WM-810 B					
WM-811 B					
WM-812 B					
WM-813 B					
WM-814 B					
WM-815 B					
WM-816 B					
WM-817 B					
WM-818 B					
WM-819 B					
WM-820 B					
WM-821 B					
WM-822 B					
WM-823 B					
WM-824 B					
WM-825 B					
WM-826 B					
WM-827 B					
WM-828 B					
WM-829 B					
WM-830 B					
WM-831 B					
WM-832 B					
WM-833 B					
WM-834 B					
WM-835 B					
WM-836 B					
WM-837 B					
WM-838 B					
WM-839 B					
WM-840 B					
WM-841 B					
WM-842 B					
WM-843 B					
WM-844 B					
WM-845 B					
WM-846 B					
WM-847 B					
WM-848 B					
WM-849 B					
WM-850 B					
WM-851 B					
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WM-856 B					
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WM-861 B					
WM-862 B					
WM-863 B					
WM-864 B					
WM-865 B					
WM-866 B					
WM-867 B					
WM-868 B					
WM-869 B					
WM-870 B					
WM-871 B					
WM-872 B					
WM-873 B					
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WM-888 B					
WM-889 B					
WM-890 B					
WM-891 B					
WM-892 B					
WM-893 B					
WM-894 B					
WM-895 B					
WM-896 B					
WM-897 B					
WM-898 B					
WM-899 B					
WM-900 B					
WM-901 B					
WM-902 B					
WM-903 B					
WM-904 B					
WM-905 B					

Figure 3 N

WM-683	E	2588	2468	2407	2723	2723	2723
WM-684	E	2590	2598	2590	2723	2723	2723
WM-685	E	2722	2723	2723	2751	2751	2751
WM-686	E	2751	2751	2751	3065	3065	3065
WM-687	E	3375	3375	3375	4128	4128	4128
WM-688	E	4131	4130	4130	4162	4162	4162
WM-689	E	4162	4162	4162	4804	4804	4804
WM-690	E	4267	4267	4267	4849	4849	4849
WM-691	E	4345	4345	4345	5988	5988	5988
WM-692	E	4485	4485	4485	6310	6310	6310
WM-693	E	4835	4835	4835	6445	6445	6445
WM-694	E	6415	6415	6415	6442	6442	6442
WM-695	E	6481	6481	6481	6482	6482	6482
WM-696	E	17923	17919	17919	10942	10942	10942
WM-697	E	17923	17920	17920	14408	14408	14408
WM-698	E	84031	84155	84155	17803	17803	17803
WM-699	E	84747	94693	94693	40301	40301	40301
WM-700	E	2281	3322	3322	78293	78293	78293
WM-701	F	3324	4172	4172	84188	84188	84188
WM-702	F	4167	4170	4170	94734	94734	94734
WM-703	F	5011	5010	5010	178344	178344	178344
WM-704	F	5009	5009	5009	2184	2184	2184
WM-705	F	6678	6678	6678	3314	3314	3314
WM-706	F	7578	7578	7578	4174	4174	4174
WM-707	F	7624	7624	7624	4718	4718	4718
WM-708	F	17935	17946	17946	5011	5011	5011
WM-709	F	45410	45399	45399	7823	7823	7823
WM-710	F	50518	50658	50658	14425	14425	14425
WM-711	F	82174	82044	82044	17845	17845	17845
WM-712	F				46301	46301	46301
WM-713	F				58844	58844	58844
WM-714	F				68902	68902	68902
WM-715	F				17841	17841	17841
WM-716	F				45385	45385	45385
WM-717	F				50514	50514	50514
WM-718	F				75836	75836	75836
WM-719	F				82092	82092	82092
WM-720	F				144070	144070	144070
WM-721	F				151883	151883	151883
WM-722	F				4109	4109	4109
WM-723	F				8876	8876	8876
WM-724	F				7580	7580	7580
WM-725	F				44158	44158	44158
WM-726	F				60463	60463	60463
WM-727	F				8443	8443	8443
WM-728	F				6432	6432	6432
WM-729	F				7844	7844	7844
WM-730	F				4837	4837	4837
WM-731	F				4480	4480	4480
WM-732	F				4849	4849	4849
WM-733	F				8443	8443	8443
WM-734	F				6432	6432	6432
WM-735	F				7844	7844	7844
WM-736	F				4837	4837	4837
WM-737	F				4480	4480	4480
WM-738	F				4849	4849	4849
WM-739	F				8443	8443	8443
WM-740	F				6432	6432	6432
WM-741	F				7844	7844	7844
WM-742	F				4837	4837	4837
WM-743	F				4480	4480	4480
WM-744	F				4849	4849	4849
WM-745	F				8443	8443	8443
WM-746	F				6432	6432	6432
WM-747	F				7844	7844	7844
WM-748	F				4837	4837	4837
WM-749	F				4480	4480	4480
WM-750	F				4849	4849	4849
WM-751	F				8443	8443	8443
WM-752	F				6432	6432	6432
WM-753	F				7844	7844	7844
WM-754	F				4837	4837	4837
WM-755	F				4480	4480	4480
WM-756	F				4849	4849	4849
WM-757	F				8443	8443	8443
WM-758	F				6432	6432	6432
WM-759	F				7844	7844	7844
WM-760	F				4837	4837	4837
WM-761	F				4480	4480	4480
WM-762	F				4849	4849	4849
WM-763	F				8443	8443	8443
WM-764	F				6432	6432	6432
WM-765	F				7844	7844	7844
WM-766	F				4837	4837	4837
WM-767	F				4480	4480	4480
WM-768	F				4849	4849	4849
WM-769	F				8443	8443	8443
WM-770	F				6432	6432	6432
WM-771	F				7844	7844	7844
WM-772	F				4837	4837	4837
WM-773	F				4480	4480	4480
WM-774	F				4849	4849	4849
WM-775	F				8443	8443	8443
WM-776	F				6432	6432	6432
WM-777	F				7844	7844	7844
WM-778	F				4837	4837	4837
WM-779	F				4480	4480	4480
WM-780	F				4849	4849	4849
WM-781	F				8443	8443	8443
WM-782	F				6432	6432	6432
WM-783	F				7844	7844	7844
WM-784	F				4837	4837	4837
WM-785	F				4480	4480	4480
WM-786	F				4849	4849	4849
WM-787	F				8443	8443	8443
WM-788	F				6432	6432	6432
WM-789	F				7844	7844	7844
WM-790	F				4837	4837	4837
WM-791	F				4480	4480	4480
WM-792	F				4849	4849	4849
WM-793	F				8443	8443	8443
WM-794	F				6432	6432	6432
WM-795	F				7844	7844	7844
WM-796	F				4837	4837	4837
WM-797	F				4480	4480	4480
WM-798	F				4849	4849	4849
WM-799	F				8443	8443	8443
WM-800	F				6432	6432	6432
WM-801	F				7844	7844	7844
WM-802	F				4837	4837	4837
WM-803	F				4480	4480	4480
WM-804	F				4849	4849	4849
WM-805	F				8443	8443	8443
WM-806	F				6432	6432	6432
WM-807	F				7844	7844	7844
WM-808	F				4837	4837	4837
WM-809	F				4480	4480	4480
WM-810	F				4849	4849	4849
WM-811	F				8443	8443	8443
WM-812	F				6432	6432	6432
WM-813	F				7844	7844	7844
WM-814	F				4837	4837	4837
WM-815	F				4480	4480	4480
WM-816	F				4849	4849	4849
WM-817	F				8443	8443	8443
WM-818	F				6432	6432	6432
WM-819	F				7844	7844	7844
WM-820	F				4837	4837	4837
WM-821	F				4480	4480	4480
WM-822	F				4849	4849	4849
WM-823	F				8443	8443	8443
WM-824	F				6432	6432	6432
WM-825	F				7844	7844	7844
WM-826	F				4837	4837	4837
WM-827	F				4480	4480	4480
WM-828	F				4849	4849	4849
WM-829	F				8443	8443	8443
WM-830	F				6432	6432	6432
WM-831	F				7844	7844	7844
WM-832	F				4837	4837	4837
WM-833	F				4480	4480	4480
WM-834	F				4849	4849	4849
WM-835	F				8443	8443	8443
WM-836	F				6432	6432	6432
WM-837	F				7844	7844	7844
WM-838	F				4837	4837	4837
WM-839	F				4480	4480	4480
WM-840	F				4849	4849	4849
WM-841	F				8443	8443	8443
WM-842	F				6432	6432	6432
WM-843	F				7844	7844	7844
WM-844	F				4837	4837	4837
WM-845	F				4480	4480	4480
WM-846	F				4849	4849	4849
WM-847	F				8443	8443	8443
WM-848	F				6432	6432	6432
WM-849	F				7844	7844	7844
WM-850	F				4837	4837	4837
WM-851	F				4480	4480	4480
WM-852	F				4849	4849	4849
WM-853	F				8443	8443	8443
WM-854	F				6432	6432	6432
WM-855	F				7844	7844	7844
WM-856	F				4837	4837	4837
WM-857	F				4480	4480	4480
WM-858	F				4849	4849	4849
WM-859	F				8443	8443	8443
WM-860	F				6432	6432	6432
WM-861	F				7844	7844	7844
WM-862	F				4837	4837	4837
WM-863	F				4480	4480	4480
WM-864	F				4849	4849	4849
WM-865	F				8443	8443	8443
WM-866	F				6432	6432	6432
WM-867	F				7844	7844	7844
WM-868	F				4837	4837	4837
WM-869	F				4480	4480	4480
WM-870	F				4849	4849	4849
WM-871	F				8443	8443	8443
WM-872	F				6432	6432	6432
WM-873	F				7844	7844	7844
WM-874	F				4837	4837	4837
WM-875	F				4480	4480	4480
WM-876	F				4849	4849	4849
WM-877	F				8443	8443	8443
WM-878	F				6432	6432	6432
WM-879	F				7844	7844	7844
WM-880	F				4837	4837	4837
WM-881	F				4480	4480	4480
WM-882	F				4849	4849	4849
WM-883	F				8443	8443	8443
WM-884</							

Figure 30.



84108
88700
162375
166886

84125
88712
162354

161441

84472

84133



WN-714 F
WN-715 F
WN-716 F
WN-717 F

Figure 4A

Rank	Normal vs Cancer	Adeno vs Normal	Squamous vs Normal	Small Cell vs Normal	Non-small Cell vs Normal	Large Cell vs Normal	Adeno vs Squamous	Adeno vs Small Cell	Squamous vs Small Cell
1	WM-61	WM-447	WM-447	WM-70	WM-341	WM-16	WM-62	WM-457	WM-276
2	WM-447	WM-652	WM-61	WM-706	WM-342	WM-26	WM-415	WM-72	WM-277
3	WM-446	WM-61	WM-277	WM-369	WM-343	WM-469	WM-152	WM-369	WM-362
4	WM-133	WM-446	WM-446	WM-447	WM-48	WM-134	WM-385	WM-78	WM-257
5	WM-119	WM-290	WM-133	WM-61	WM-340	WM-847	WM-347	WM-79	WM-363
6	WM-278	WM-363	WM-134	WM-652	WM-346	WM-277	WM-134	WM-73	WM-347
7	WM-134	WM-133	WM-363	WM-282	WM-47	WM-310	WM-36	WM-64	WM-53
8	WM-383	WM-341	WM-362	WM-341	WM-339	WM-446	WM-108	WM-320	WM-254
9	WM-282	WM-285	WM-276	WM-456	WM-389	WM-446	WM-99	WM-419	WM-17
10	WM-362	WM-366	WM-706	WM-134	WM-569	WM-221	WM-151	WM-85	WM-252
11	WM-120	WM-282	WM-203	WM-203	WM-447	WM-648	WM-289	WM-82	WM-431
12	WM-290	WM-302	WM-466	WM-646	WM-652	WM-557	WM-363	WM-53	WM-513
13	WM-65	WM-310	WM-366	WM-455	WM-154	WM-290	WM-61	WM-412	WM-446
14	WM-277	WM-292	WM-85	WM-65	WM-587	WM-328	WM-117	WM-440	WM-355
15	WM-70	WM-120	WM-70	WM-685	WM-456	WM-447	WM-211	WM-455	WM-447
16	WM-369	WM-134	WM-341	WM-473	WM-450	WM-084	WM-362	WM-313	WM-133
17	WM-17	WM-276	WM-429	WM-343	WM-283	WM-183	WM-133	WM-456	WM-245
18	WM-473	WM-428	WM-347	WM-466	WM-207	WM-190	WM-414	WM-86	WM-52
19	WM-47	WM-277	WM-17	WM-341	WM-436	WM-586	WM-277	WM-70	WM-96
20	WM-203	WM-20	WM-47	WM-340	WM-384	WM-397	WM-141	WM-238	WM-238
21	WM-276	WM-119	WM-431	WM-363	WM-61	WM-466	WM-64	WM-246	WM-243
22	WM-279	WM-340	WM-62	WM-339	WM-167	WM-20	WM-135	WM-190	WM-138
23	WM-62	WM-48	WM-473	WM-457	WM-382	WM-17	WM-447	WM-418	WM-62
24	WM-366	WM-399	WM-384	WM-86	WM-285	WM-545	WM-383	WM-83	WM-580
25	WM-456	WM-450	WM-438	WM-506	WM-550	WM-47	WM-338	WM-257	WM-134
26	WM-428	WM-47	WM-452	WM-72	WM-203	WM-191	WM-63	WM-138	WM-240
27	WM-384	WM-343	WM-282	WM-287	WM-119	WM-147	WM-142	WM-47	WM-256
28	WM-287	WM-17	WM-369	WM-82	WM-282	WM-480	WM-446	WM-252	WM-203
29	WM-420	WM-583	WM-290	WM-528	WM-866	WM-590	WM-186	WM-282	WM-111
30	WM-292	WM-70	WM-278	WM-85	WM-383	WM-218	WM-111	WM-60	WM-95
31	WM-431	WM-706	WM-456	WM-73	WM-429	WM-285	WM-445	WM-88	WM-247
32	WM-455	WM-346	WM-673	WM-138	WM-11	WM-652	WM-455	WM-325	WM-157
33	WM-20	WM-466	WM-340	WM-384	WM-200	WM-651	WM-276	WM-402	WM-242
34	WM-340	WM-646	WM-55	WM-83	WM-451	WM-366	WM-444	WM-411	WM-556
35	WM-19	WM-394	WM-455	WM-450	WM-473	WM-403	WM-181	WM-405	WM-63
36	WM-389	WM-336	WM-645	WM-310	WM-220	WM-418	WM-35	WM-75	WM-239
37	WM-63	WM-294	WM-138	WM-277	WM-685	WM-430	WM-205	WM-417	WM-234
38	WM-438	WM-339	WM-420	WM-79	WM-338	WM-456	WM-456	WM-387	WM-274

Figure 4B

39	WM-450	WM-473	WM-150	WM-207	WM-71	WM-714	WM-39	WM-26	WM-370
40	WM-468	WM-369	WM-369	WM-278	WM-268	WM-546	WM-82	WM-410	WM-301
41	WM-296	WM-38	WM-279	WM-290	WM-70	WM-109	WM-17	WM-420	WM-449
42	WM-343	WM-283	WM-342	WM-366	WM-545	WM-302	WM-203	WM-184	WM-74
43	WM-341	WM-685	WM-471	WM-472	WM-575	WM-567	WM-83	WM-567	WM-261
44	WM-339	WM-66	WM-574	WM-420	WM-446	WM-375	WM-412	WM-66	WM-467
45	WM-55	WM-120	WM-120	WM-147	WM-120	WM-131	WM-96	WM-391	WM-237
46	WM-66	WM-55	WM-20	WM-55	WM-267	WM-706	WM-74	WM-340	WM-262
47	WM-48	WM-307	WM-287	WM-669	WM-466	WM-398	WM-457	WM-428	WM-295
48	WM-38	WM-278	WM-83	WM-357	WM-347	WM-359	WM-431	WM-198	WM-288
49	WM-138	WM-342	WM-154	WM-429	WM-153	WM-55	WM-340	WM-312	WM-384
50	WM-310	WM-429	WM-128	WM-279	WM-38	WM-488	WM-48	WM-152	WM-37

2803	5927	10300
SAA 42-67 (2802.1)	SAA 32-85 (5925.3)	SAA 6-97 (10299.1)
3168	6874	10866
SAA 69-97 (3167.3)	SAA 26-88 (6873.3)	SAA 4-101 (10871.8)
3277	7776	10851
SAA 39-68 (3276.6)	SAA 1-68 (7774.6)	SAA 5-102 (10853.7)
3552	7941	
SAA 38-70 (3552)	SAA 18-88 (7939.5)	
3897	8152	
SAA 64-98 (3897.2)	SAA 25-98 (8150)	
4300	8952	
SAA 54-93 (4302.5)	SAA 6-85 (8950)	
4490	9233	
SAA 53-93 (4489)	SAA 16-97 (9235)	
4655		
SAA 5-44 (4655.0)		

Figure 5

Figure 5

Figure 6

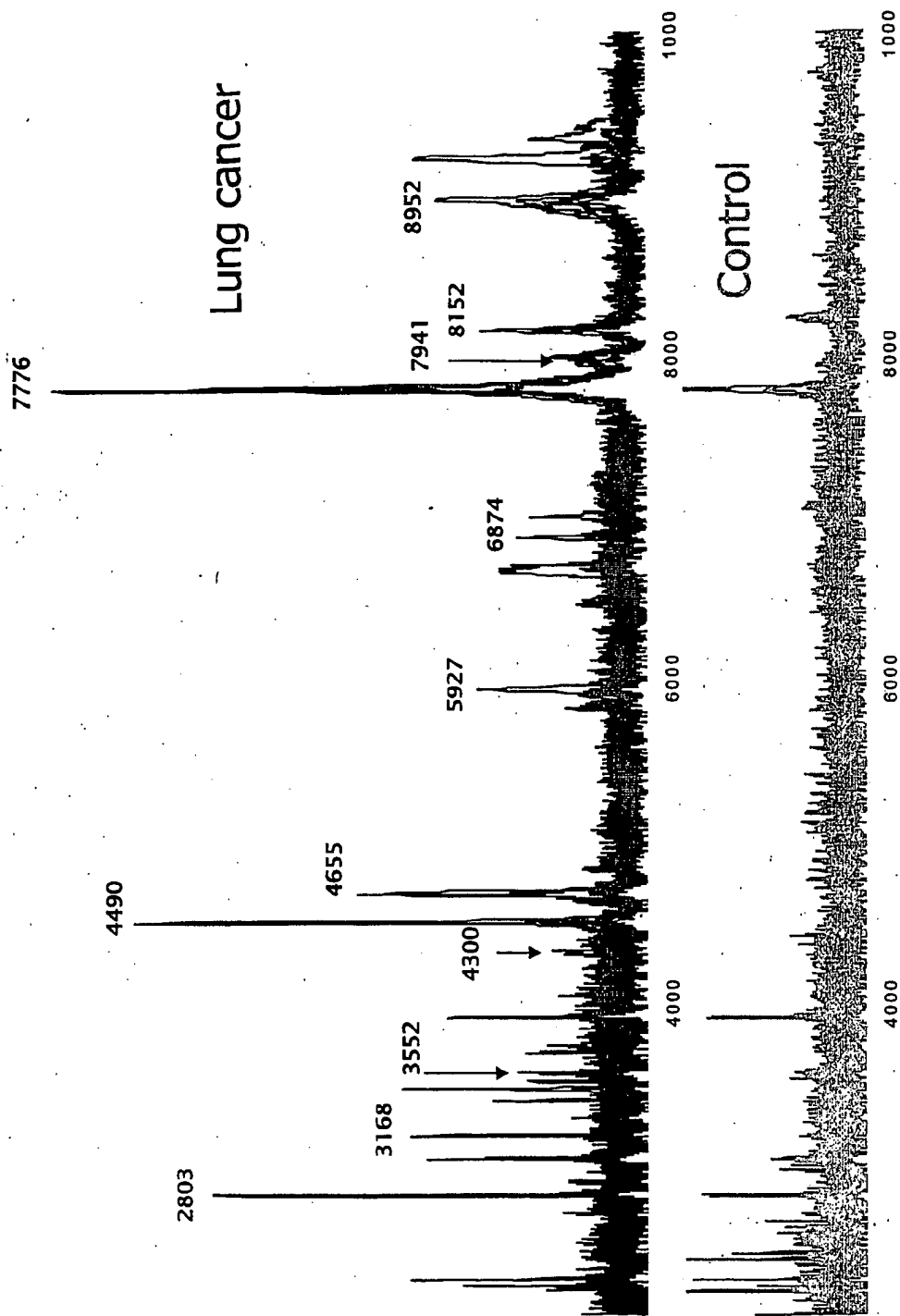


FIGURE 7
Protein Profile of Selected Samples Q Fraction 1 WCX2

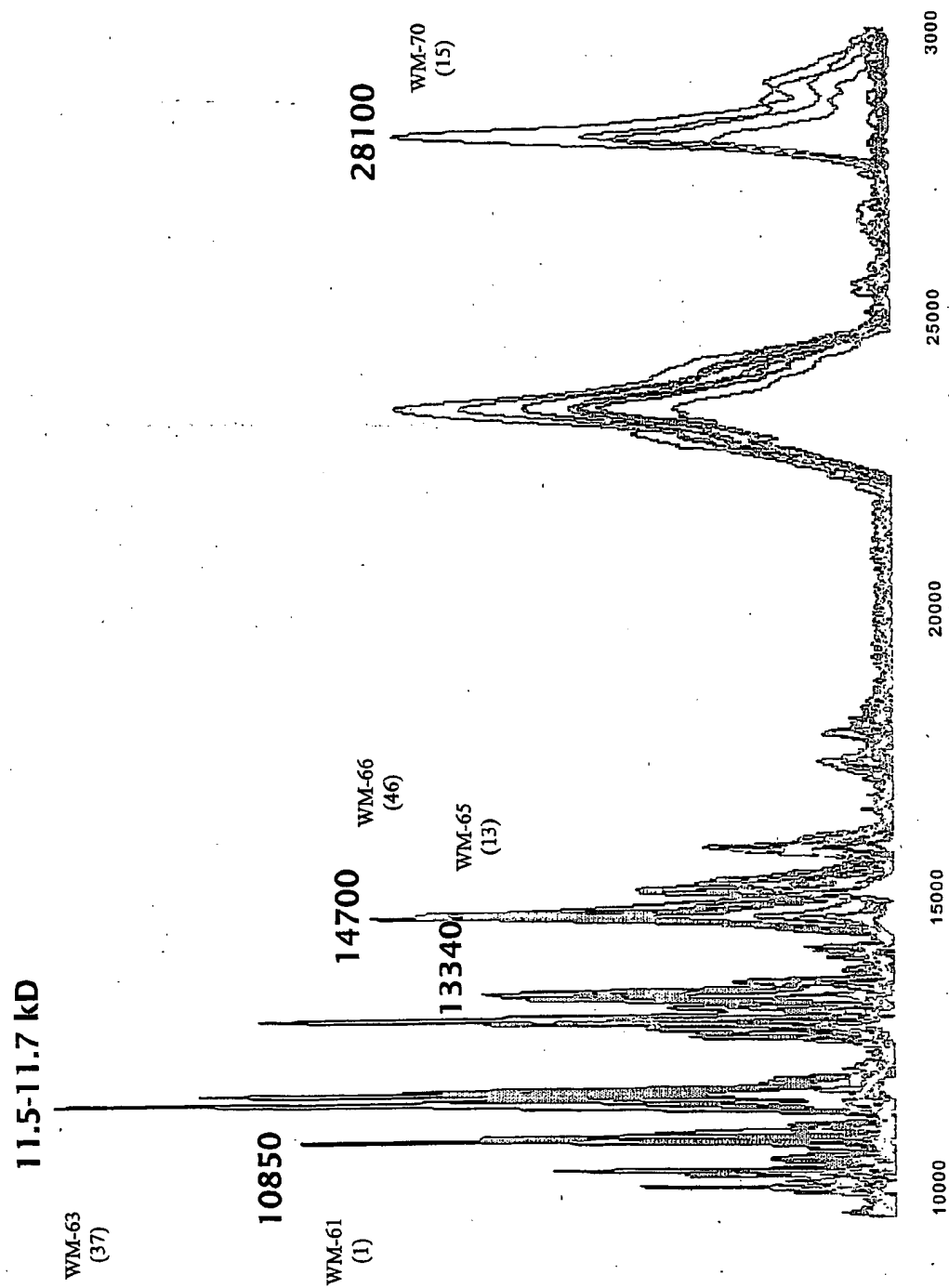


Figure 8
Protein Profile of Selected Samples
Q Fraction 1 WCX2

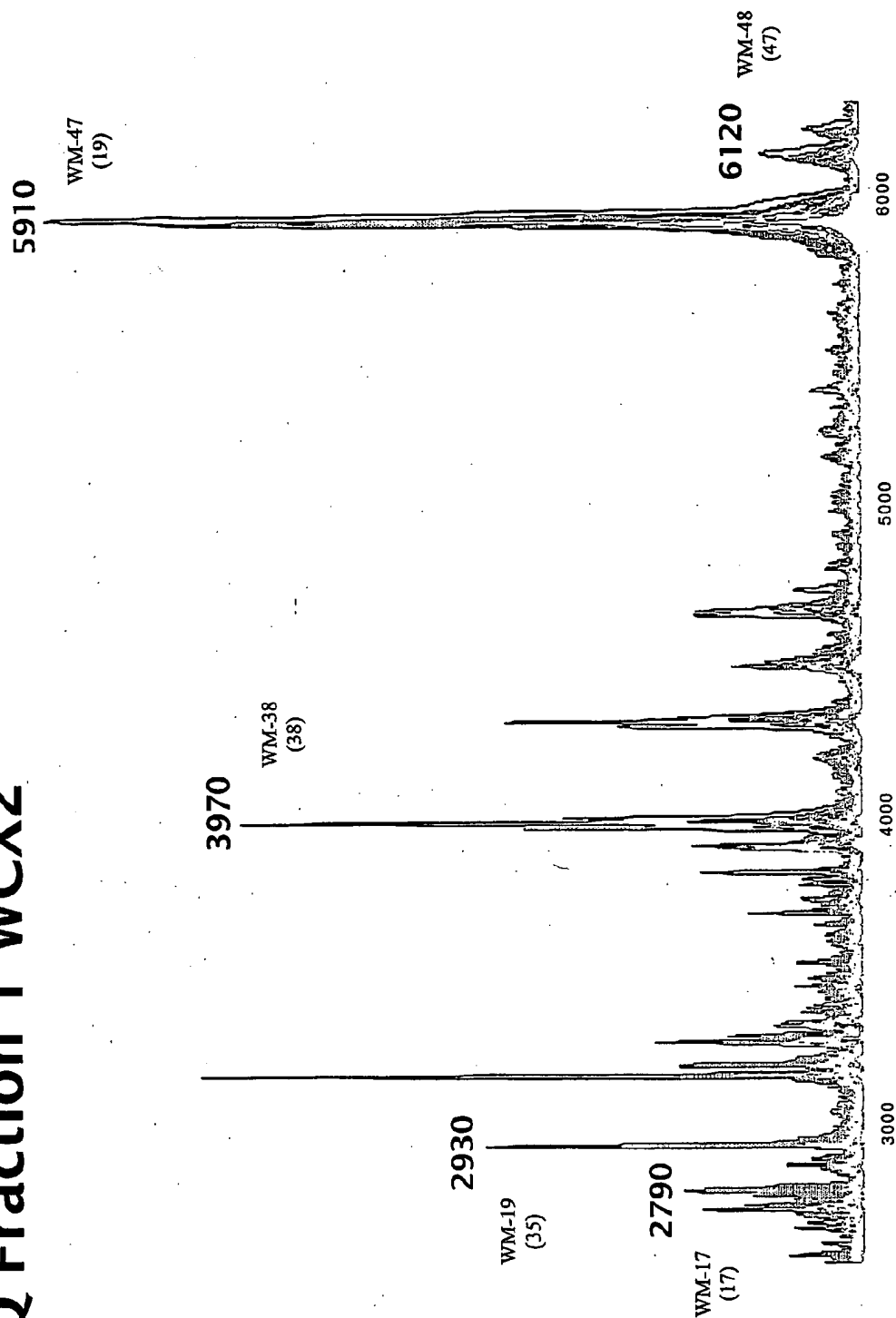


Figure 9
Protein Profile of Selected Samples
Q Fraction 2 WCX2

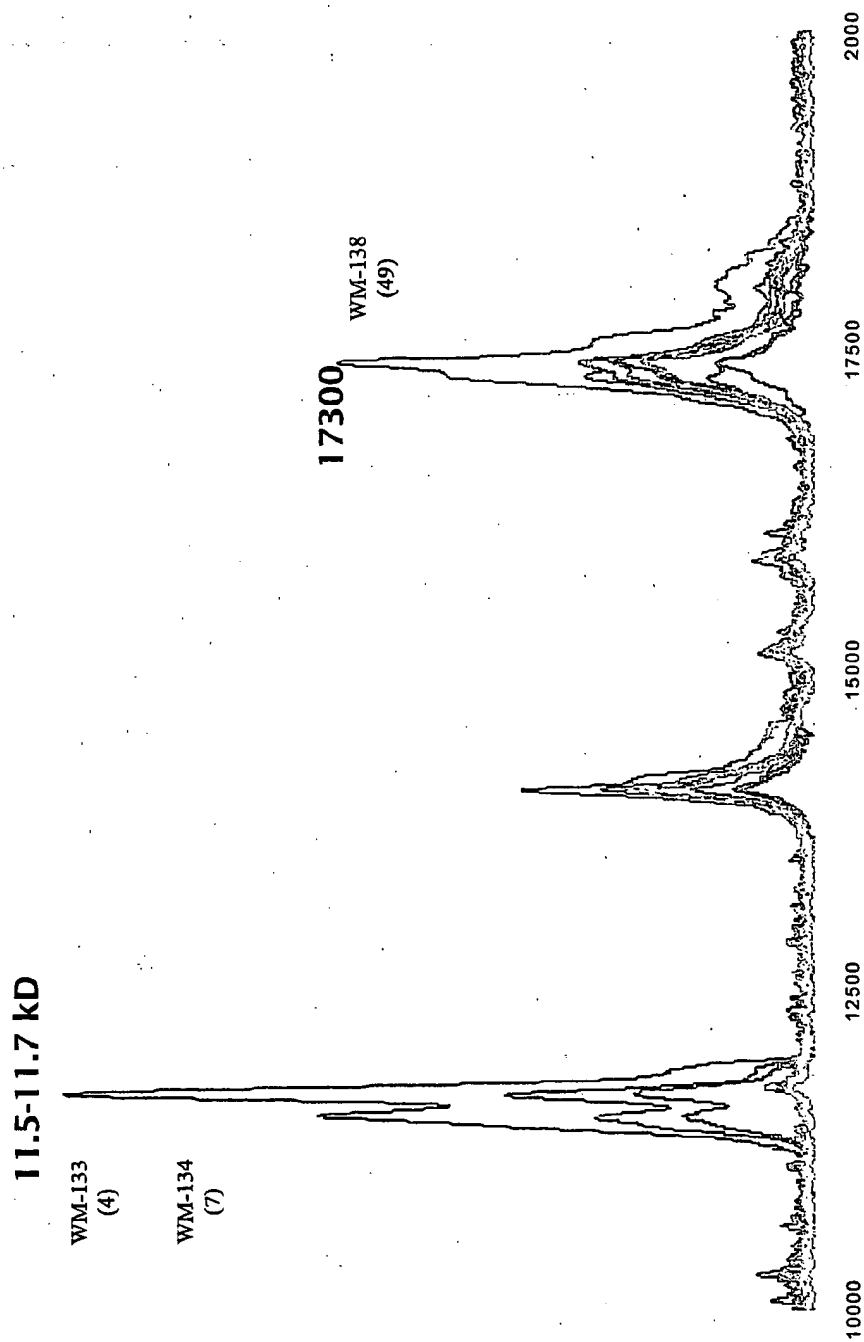


Figure 10
Protein Profile of Selected Samples
Q Fraction 2 WCX2

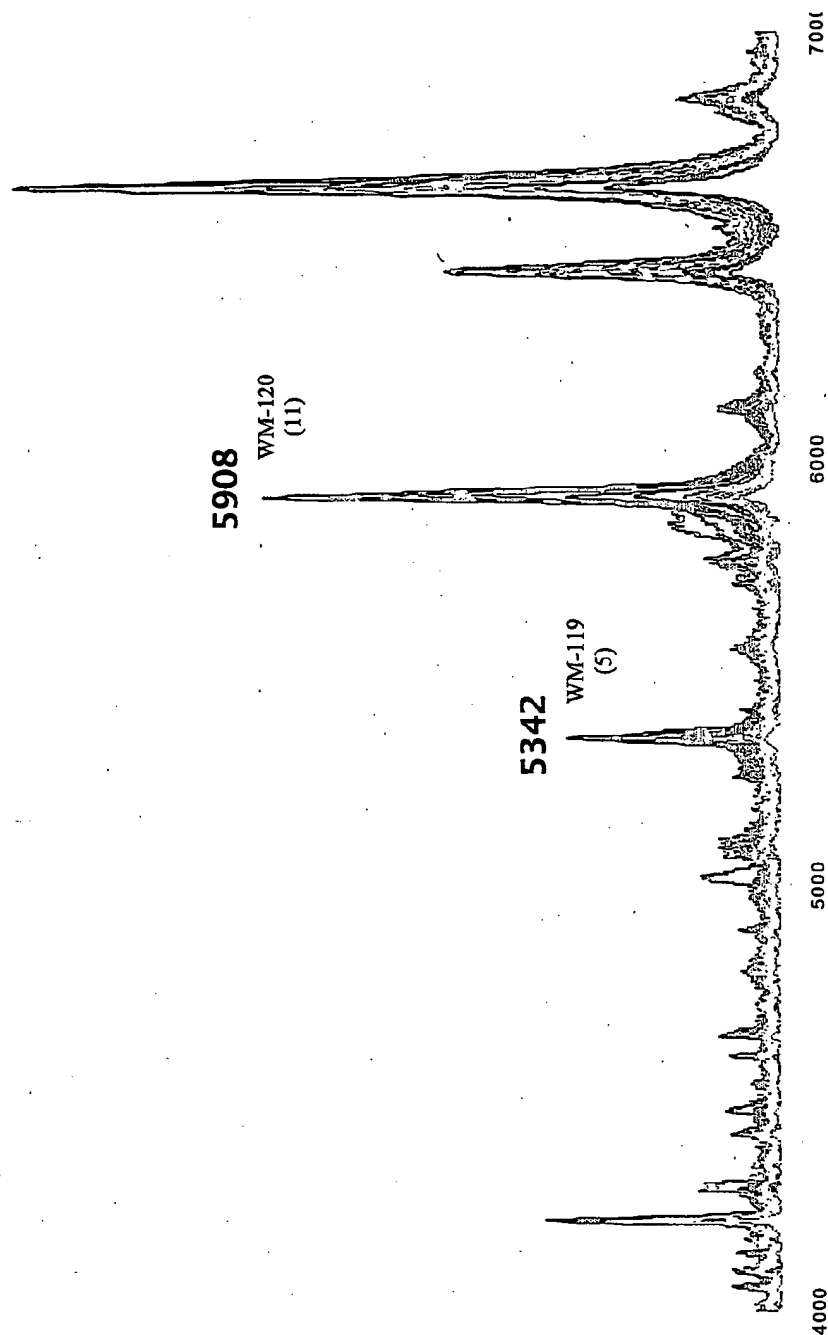


Figure 11
Protein Profile of Selected Samples
Q Fraction 4 WCX2

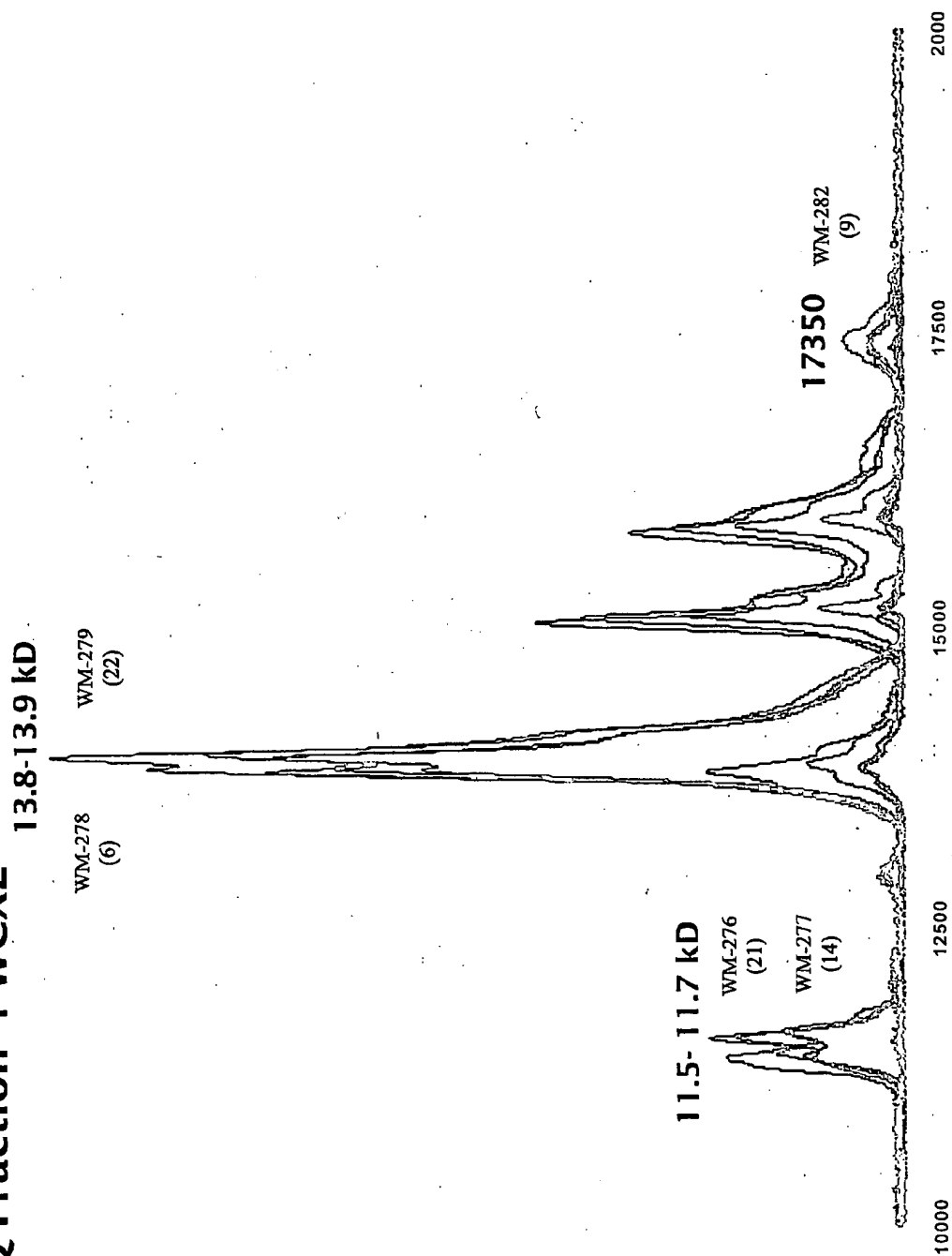


Figure 12
Protein Profile of Selected Samples
Q Fraction 4 WCX2

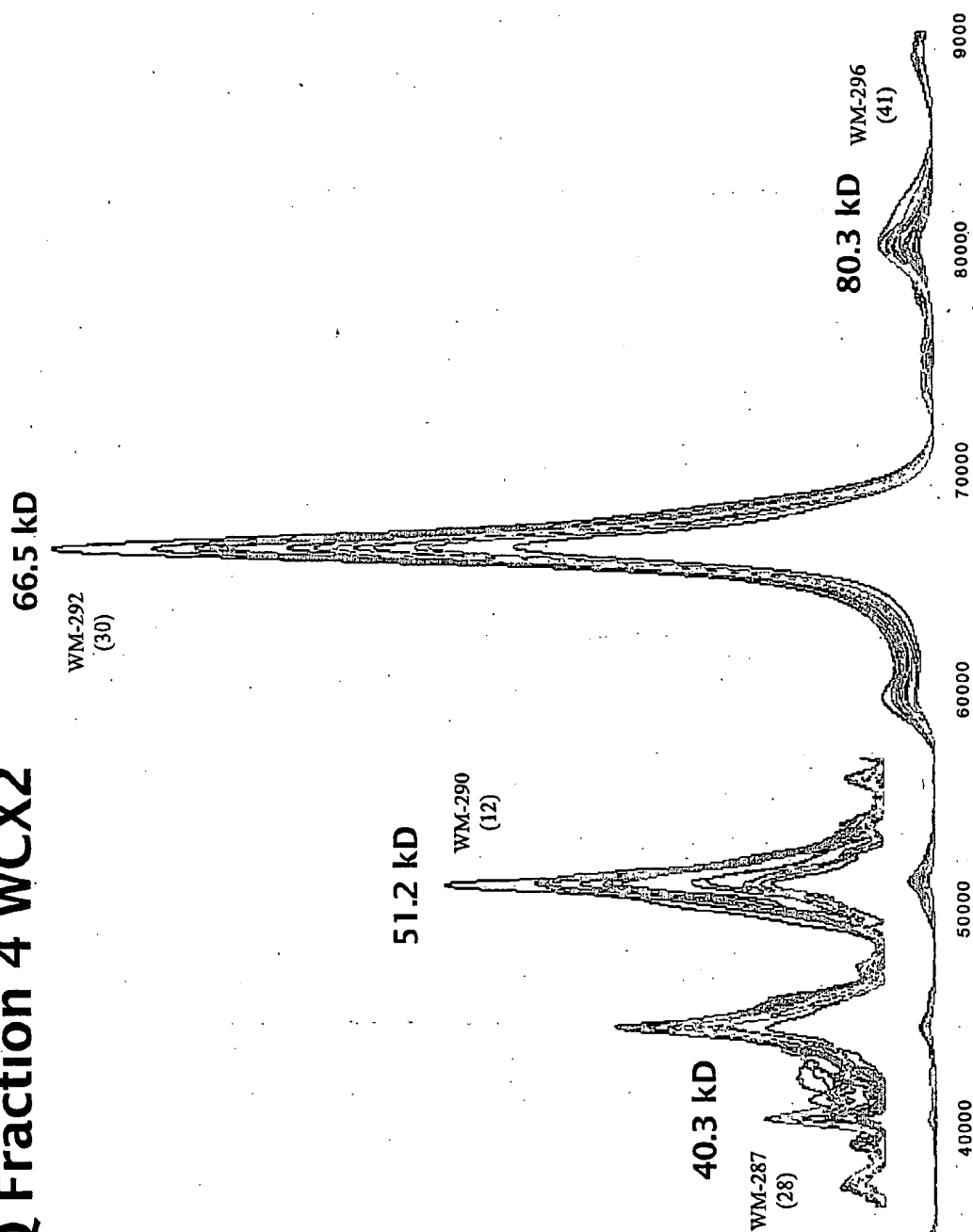


Figure 13
Protein Profile of Selected Samples
Q Fraction 5 WCX2

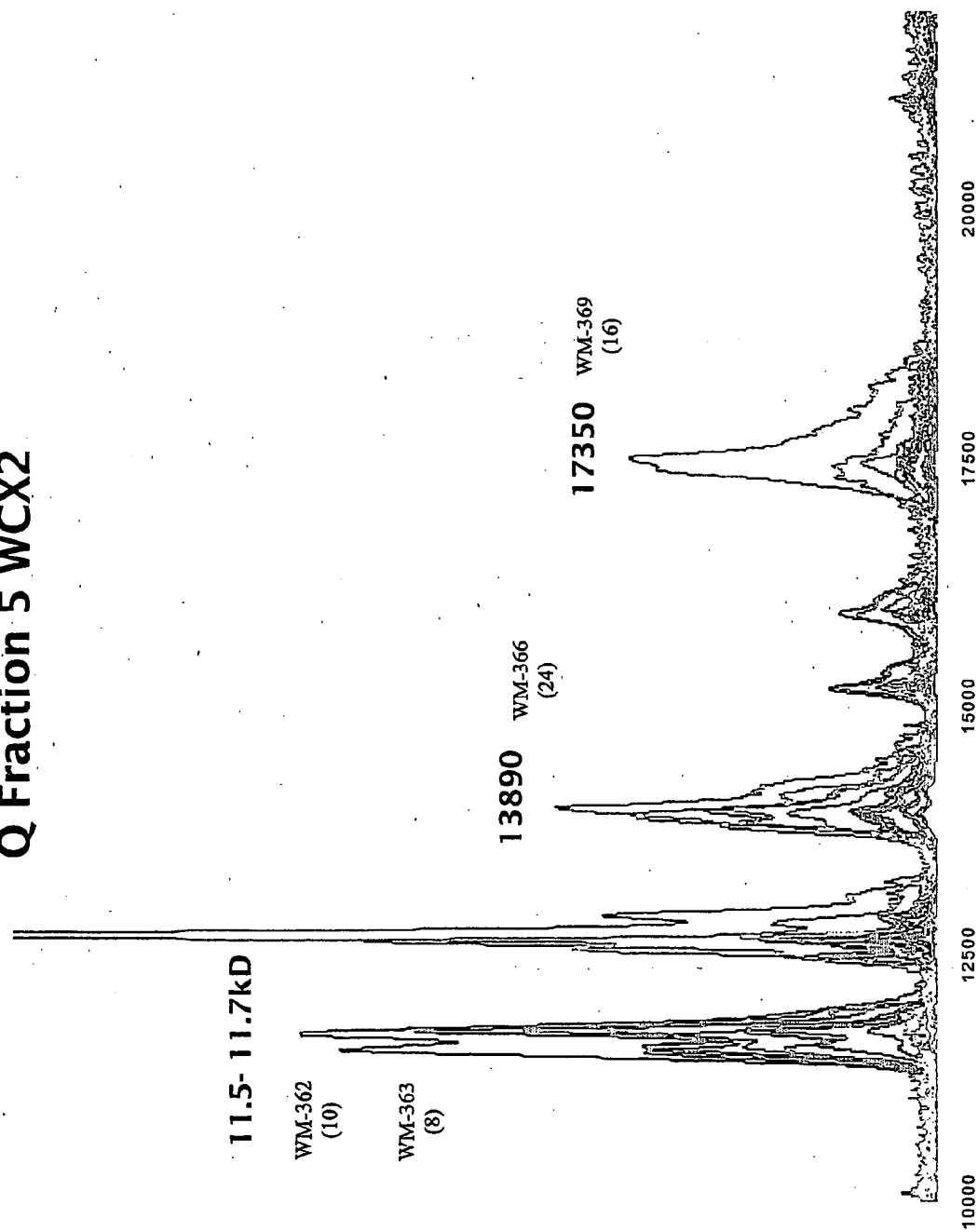


Figure 14
Protein Profile of Selected Samples
Q Fraction 5 WCX2

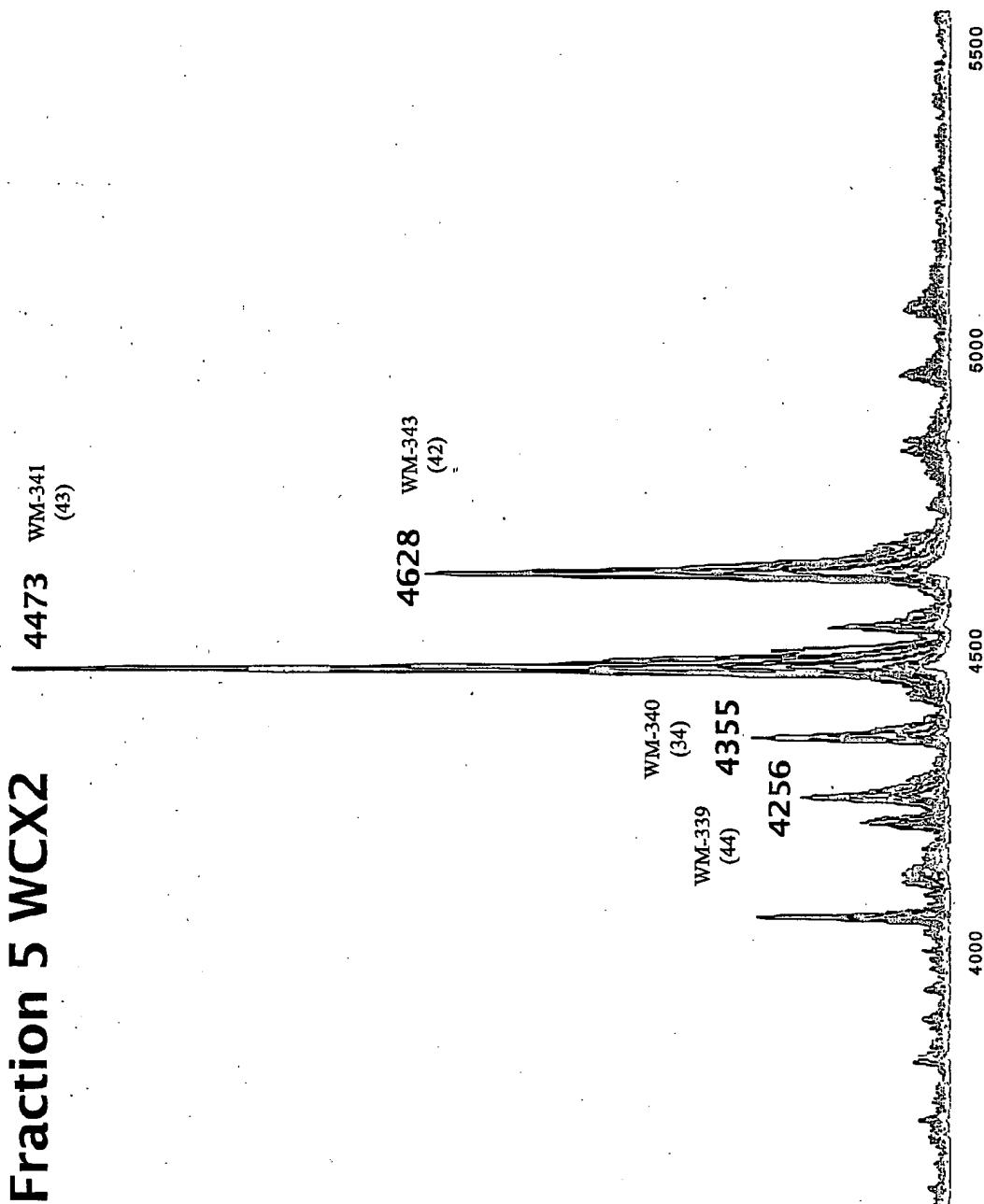


Figure 15
Protein Profile of Selected Samples
Q Fraction 6 WCX2

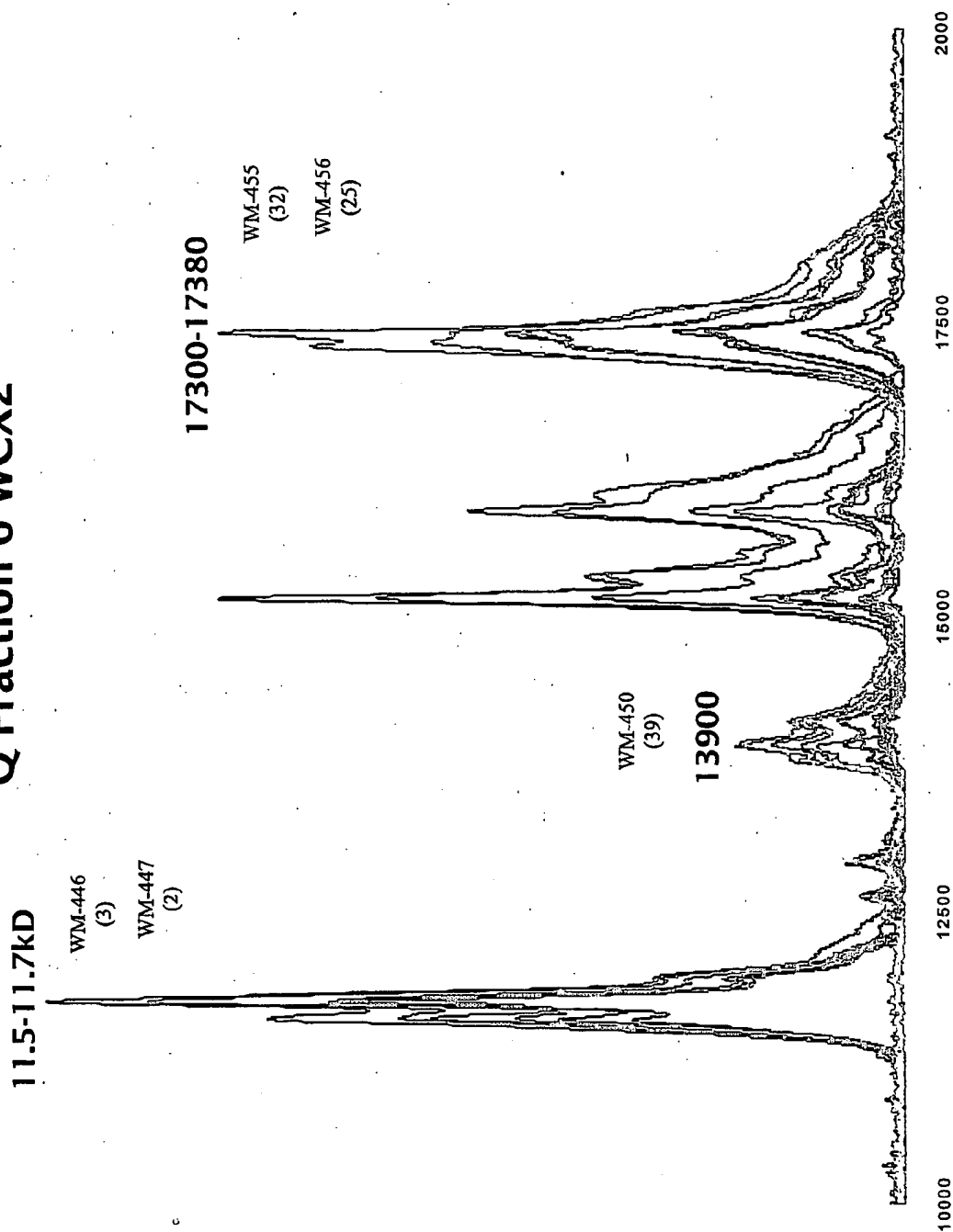


Figure 16
Protein Profile of Selected Samples
Q Fraction 6 WCX2

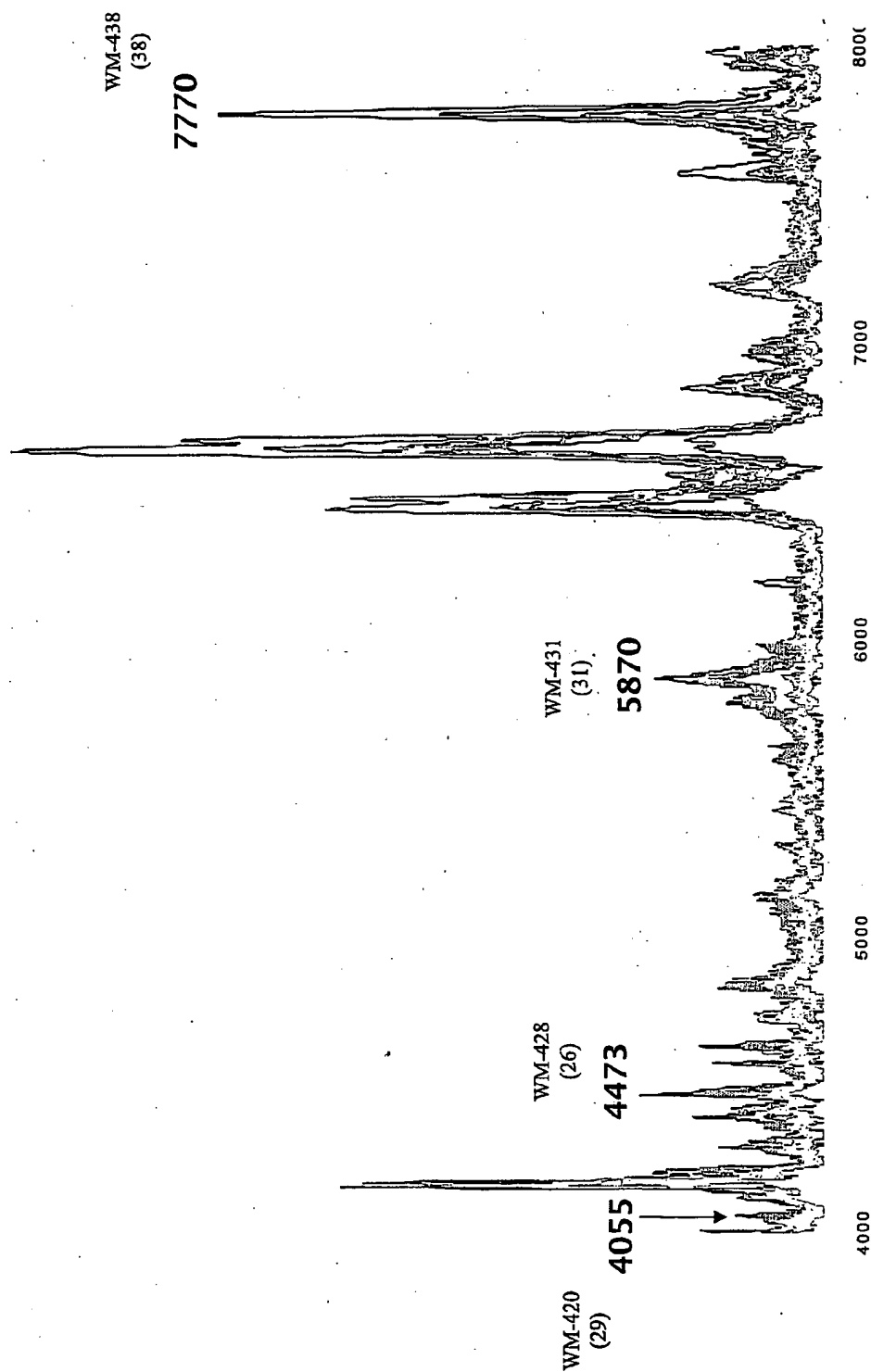


Figure 17
Protein Profile of Selected Samples
Q Fraction 2 IMAC-Cu(II)

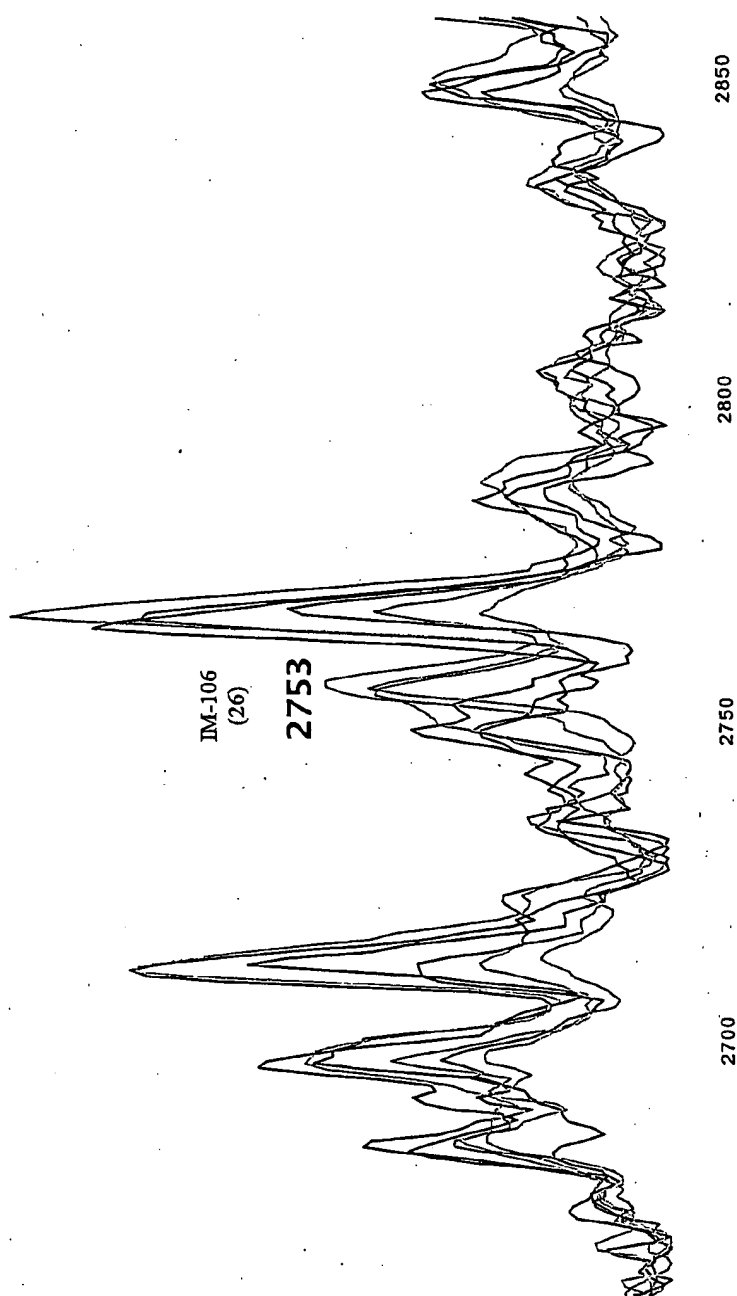


Figure 18
Protein Profile of Selected Samples
Q Fraction 2 IMAC-Cu(II)

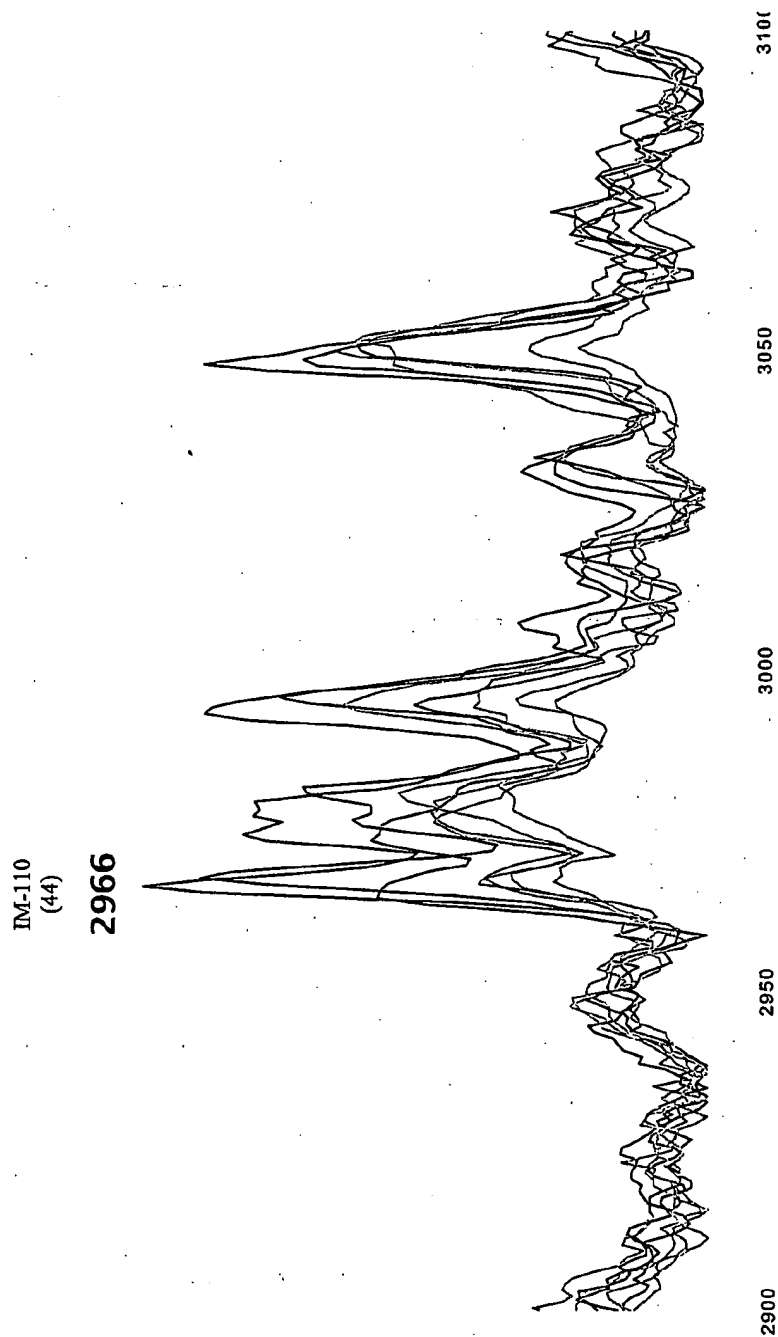


Figure 19
Protein Profile of Selected Samples
Q Fraction 2 IMAC-Cu(II)

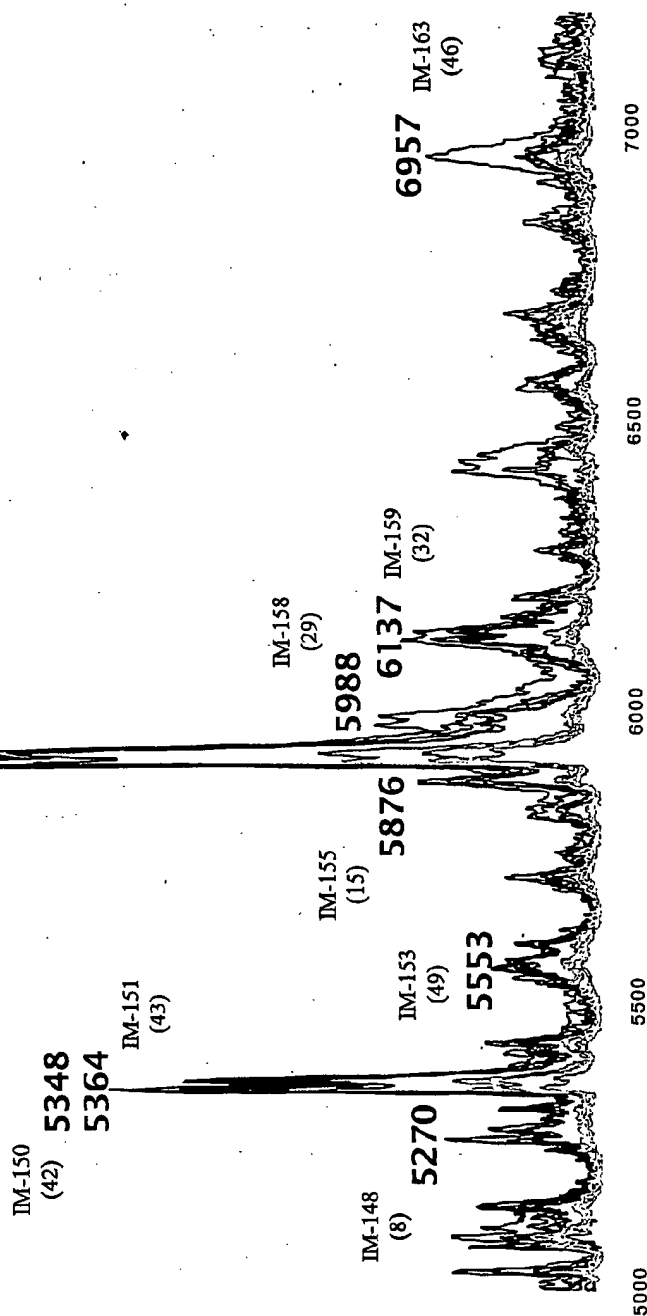


Figure 20
Protein Profile of Selected Samples
Q Fraction 2 IMAC-Cu(II)

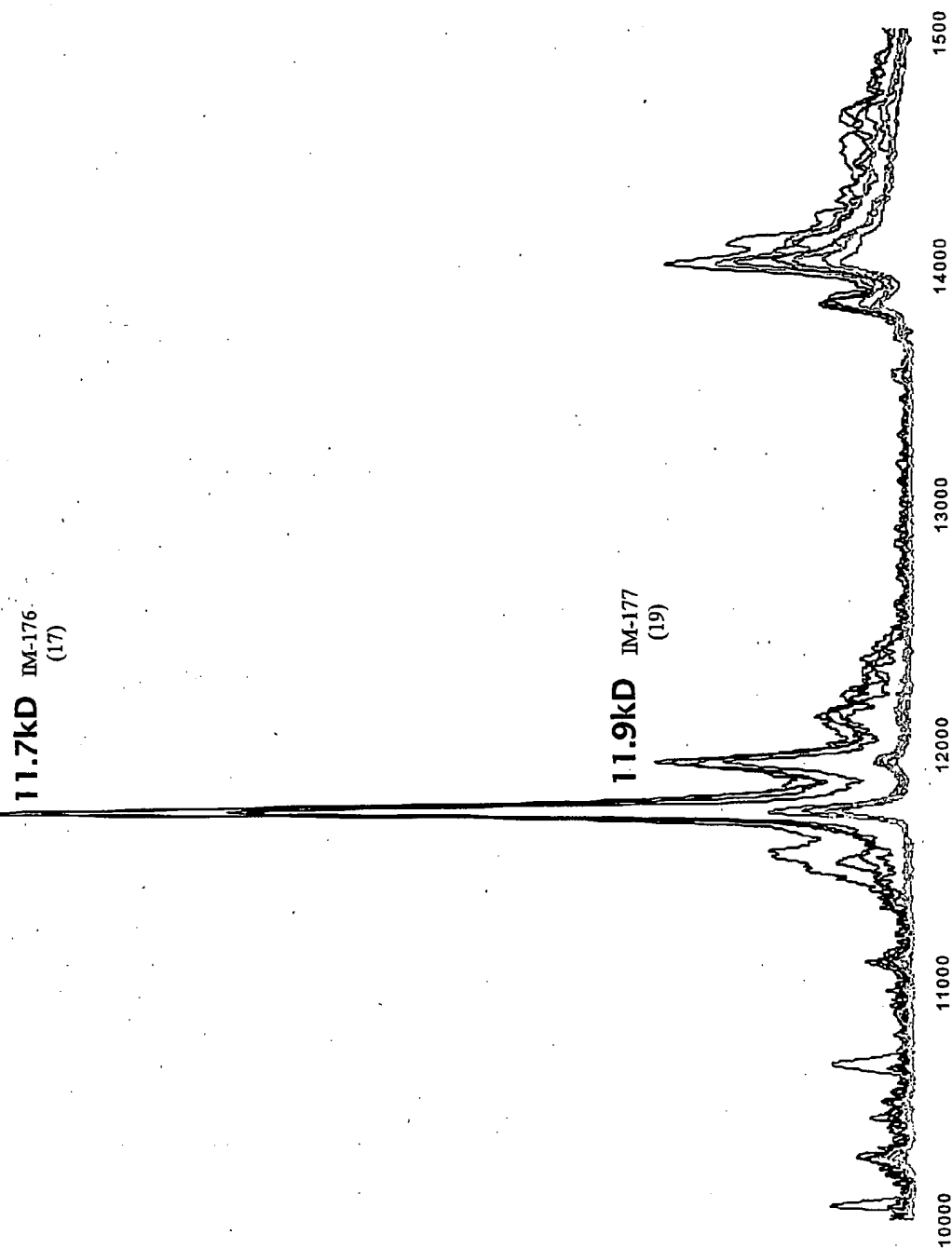


Figure 21
Protein Profile of Selected Samples
Q Fraction 3 IMAC-Cu(II)

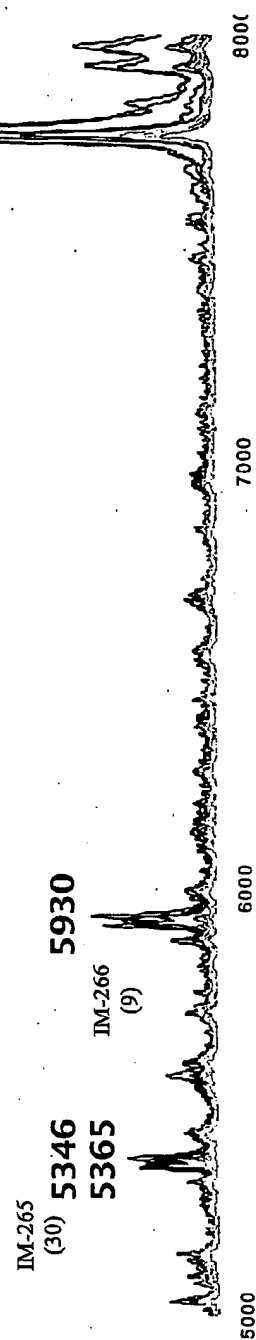


Figure 22
Protein Profile of Selected Samples
Q Fraction 3 IMAC-Cu(II)

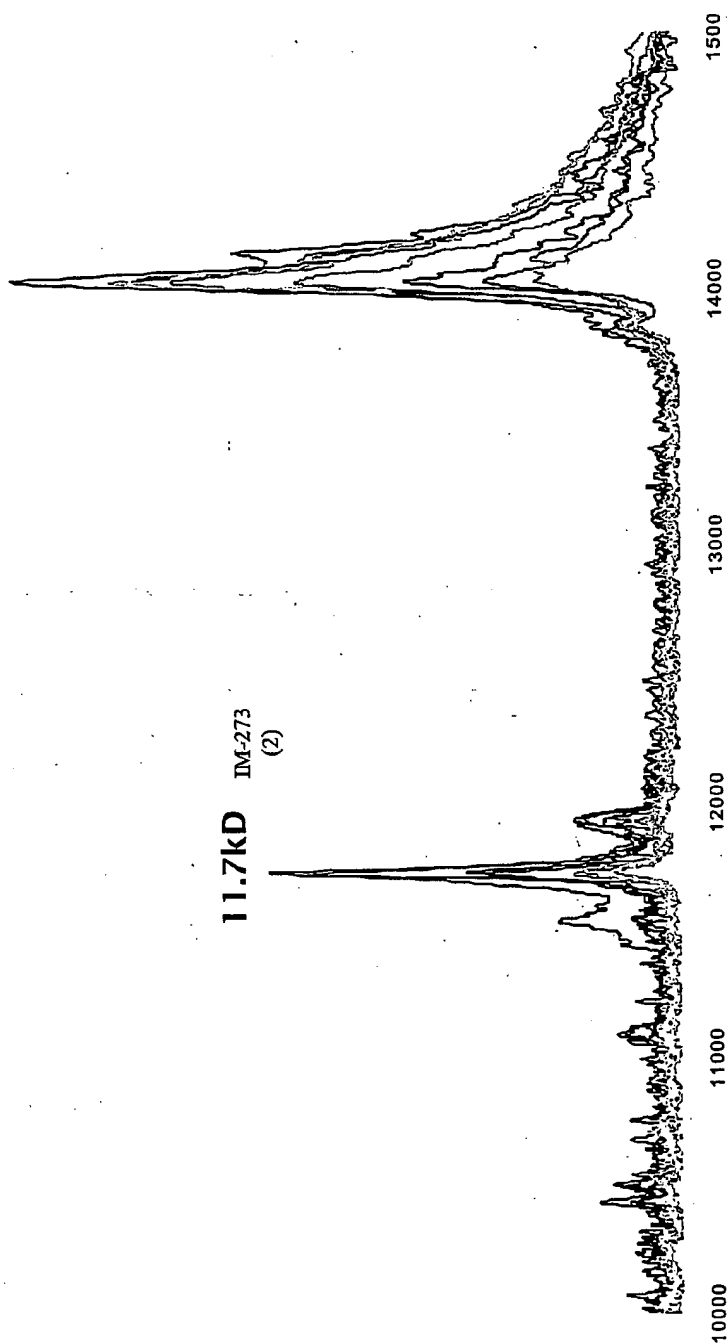


Figure 23
Protein Profile of Selected Samples
Q Fraction 5 IMAC-Cu(II)

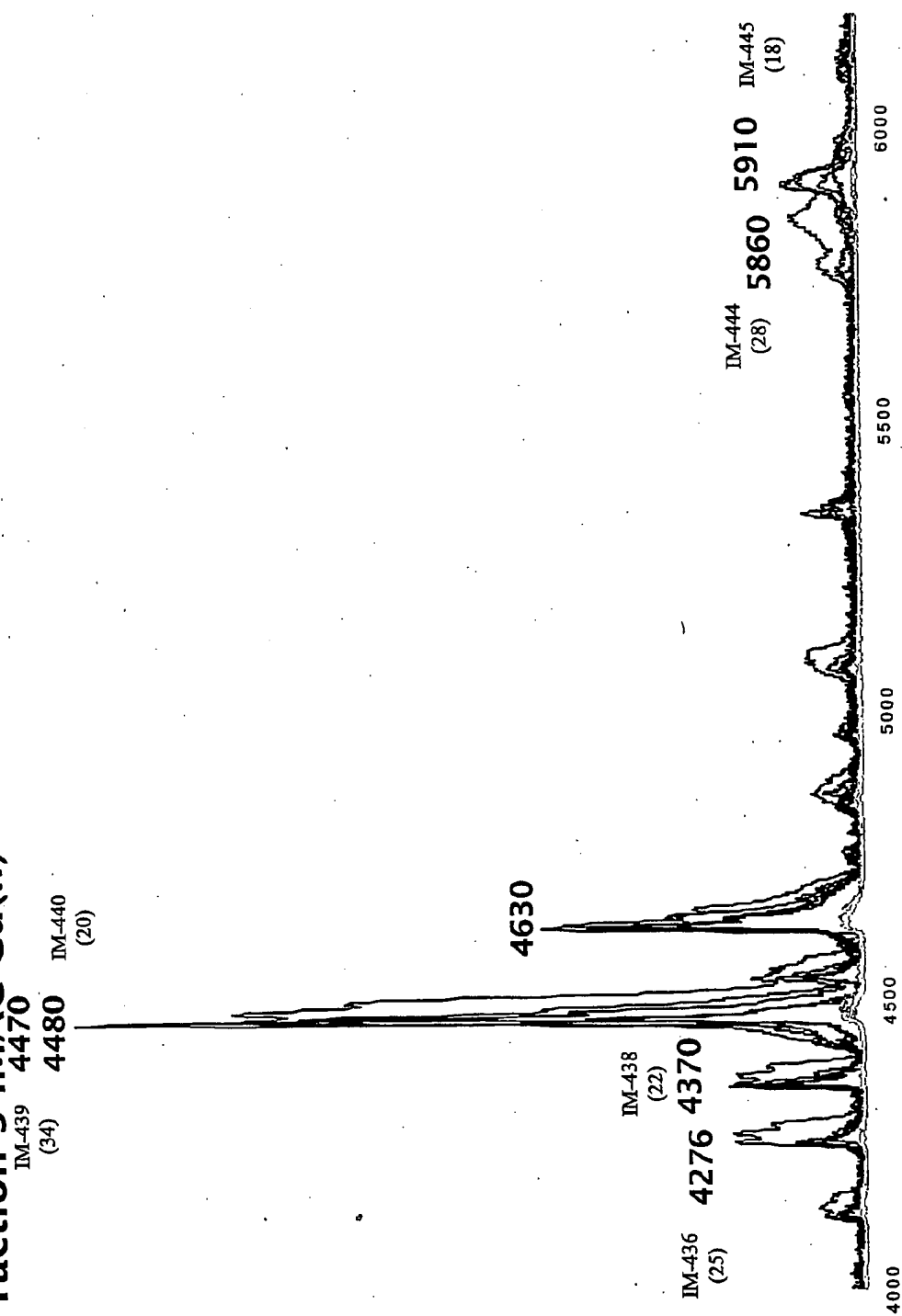


Figure 24
Protein Profile of Selected Samples
Q Fraction 5 IMAC-Cu(II)

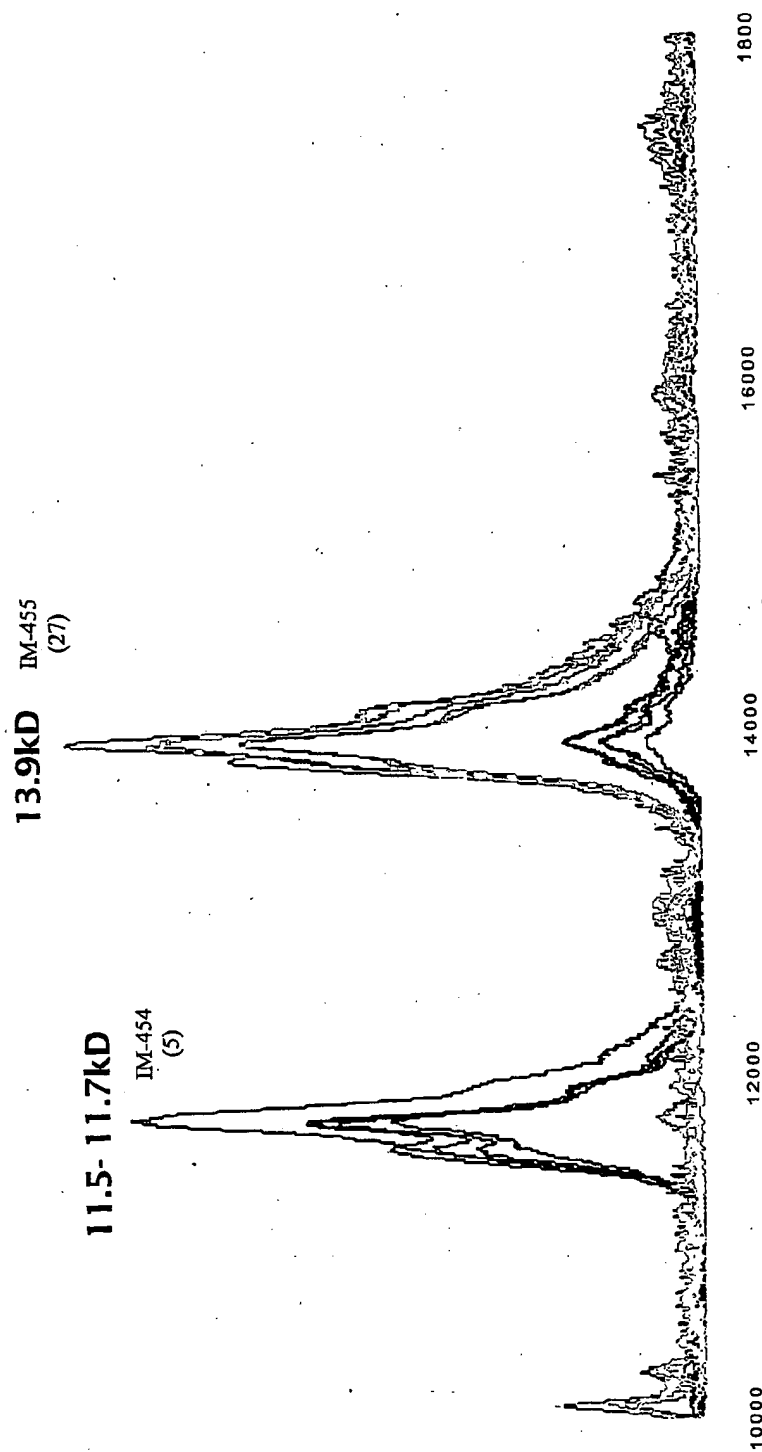


Figure 25
Protein Profile of Selected Samples
Q Fraction 5 IMAC-Cu(II)

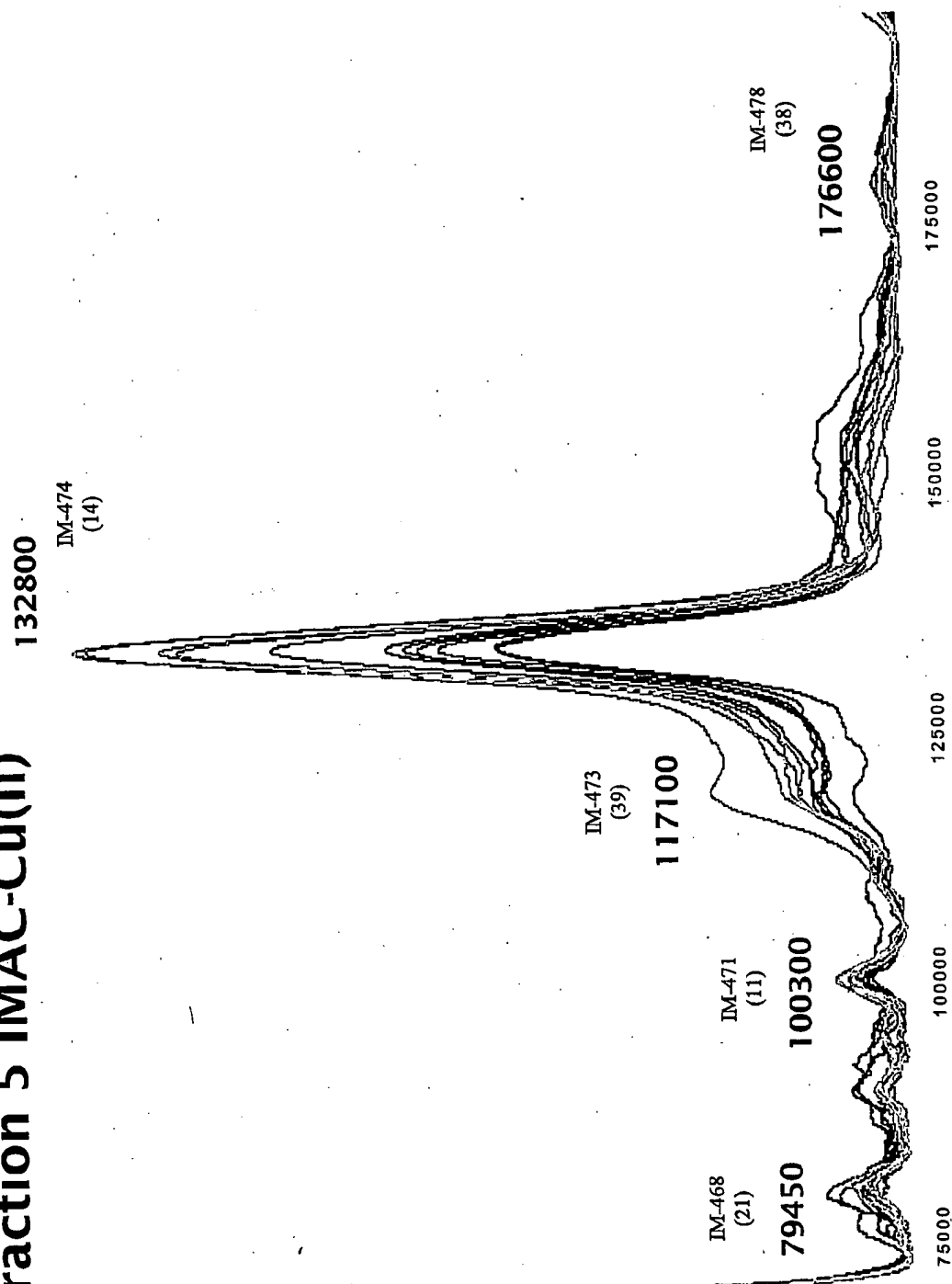


Figure 26
Protein Profile of Selected Samples
Q Fraction 6 IMAC-Cu(II)

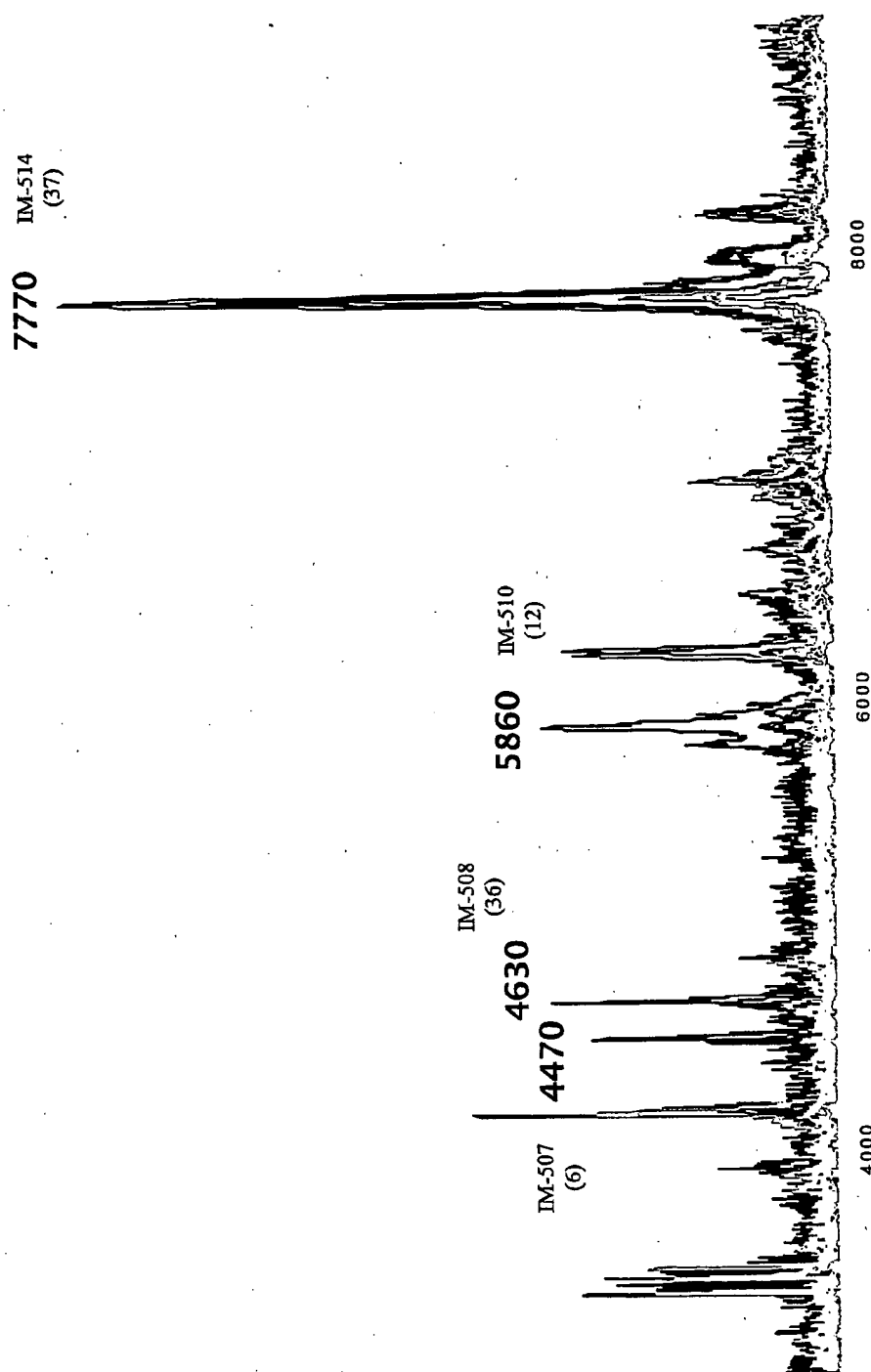


Figure 27
Protein Profile of Selected Samples
Q Fraction 6 IMAC-Cu(II)

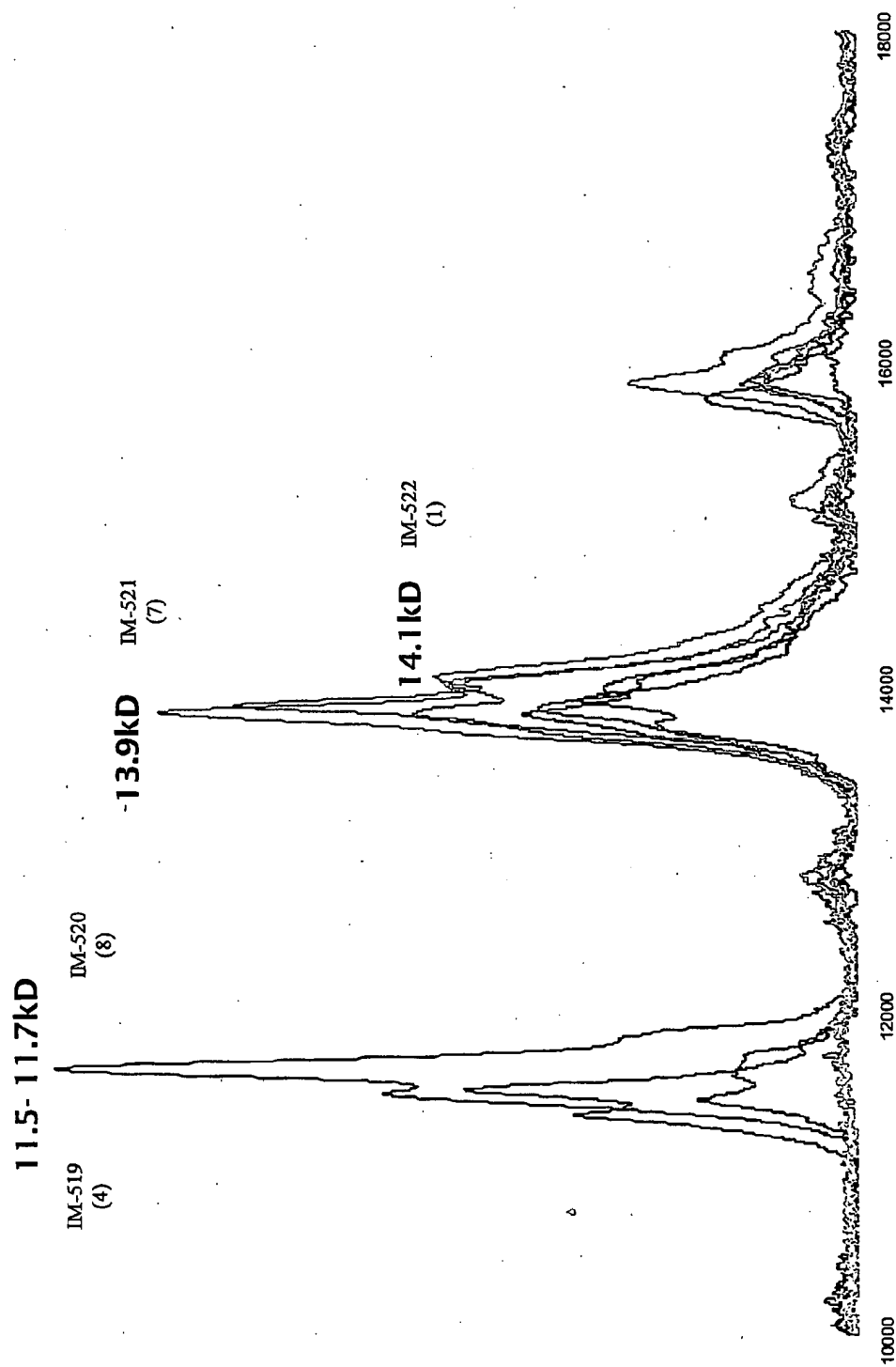
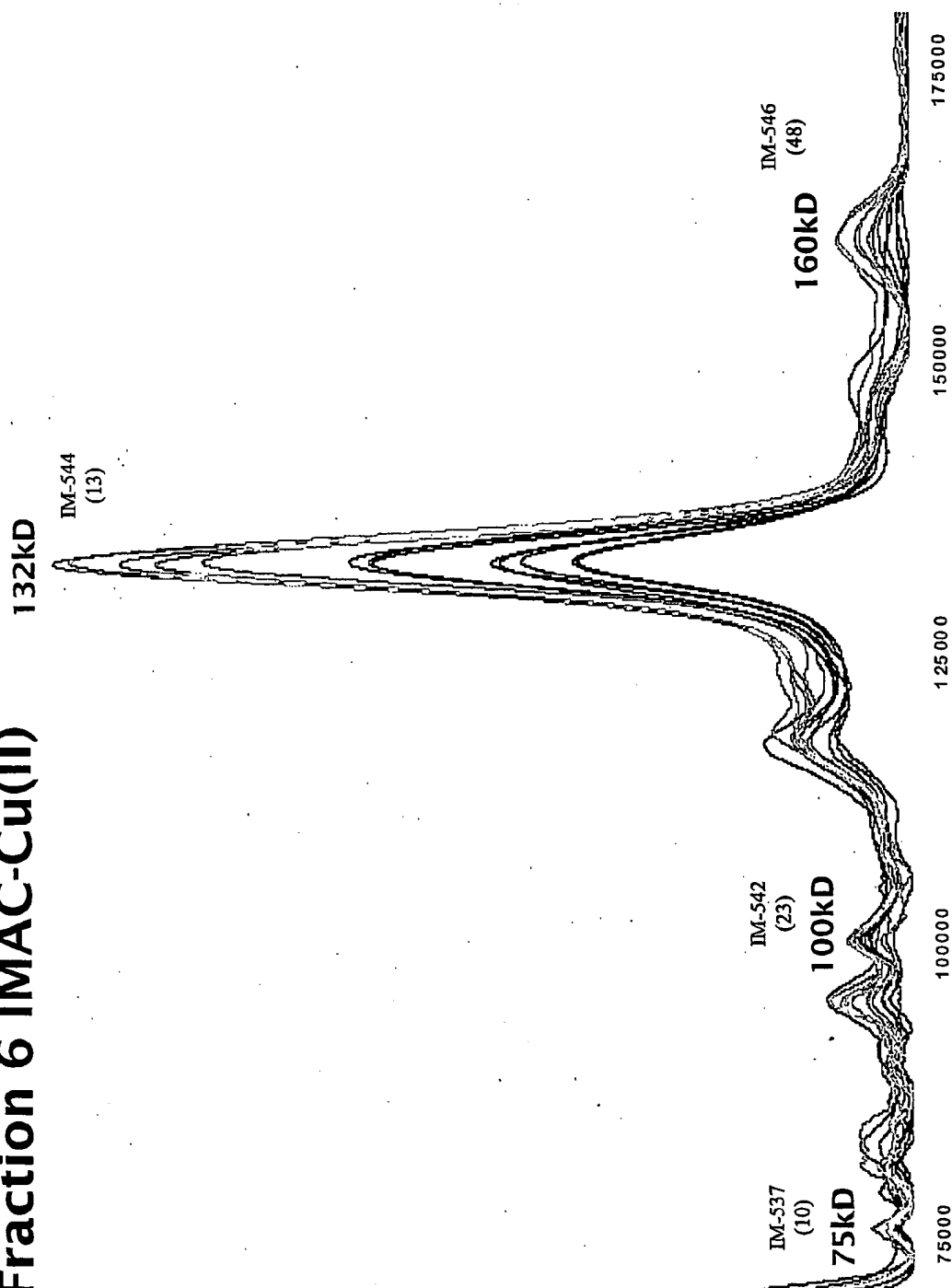


Figure 28
Protein Profile of Selected Samples
Q Fraction 6 IMAC-Cu(II)



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